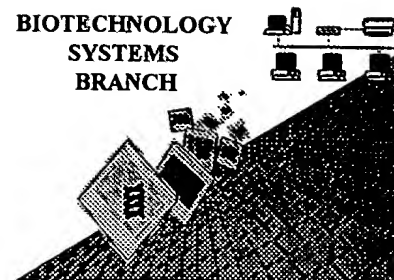


*Signature*

# **RAW SEQUENCE LISTING** **ERROR REPORT**



**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:**

Application Serial Number: 08/836,734  
Art Unit / Team No. : 1634  
Date Processed by STIC: 6/16/98

**THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.**

**PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:**

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**ARTI SHAH 703-308-4212**

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:36:21

INPUT SET: S26725.raw

This Raw Listing contains the General  
Information Section and those Sequences  
containing ERRORS.

## SEQUENCE LISTING

## (1) General Information:

(i) APPLICANT: BECKMANN, JACQUES  
RICHARD, ISABELLE

(ii) TITLE OF INVENTION: LGMD GENE CODING FOR A CALCIUM DEPENDENT PROTEASE

(iii) NUMBER OF SEQUENCES: 67

## (iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: NIXON &amp; VANDERHYE P.C.

(B) STREET: 1100 NORTH GLEBE ROAD

(C) CITY: ARLINGTON

(D) STATE: VIRGINIA

(E) COUNTRY: U.S.A.

(F) ZIP: 22201-4714

## (v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk

(B) COMPUTER: IBM PC compatible

(C) OPERATING SYSTEM: PC-DOS/MS-DOS

(D) SOFTWARE: PatentIn Release #1.0, Version #1.30

## (vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: US 08/836,734

(B) FILING DATE: 02-JUL-1997

(C) CLASSIFICATION:

## (vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: EP 94402668.1

(B) FILING DATE: 22-NOV-1994

## (viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: WILSON, MARY J.

(B) REGISTRATION NUMBER: 32,955

(C) REFERENCE/DOCKET NUMBER: 960-29

## (ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: (703) 816-4000

(B) TELEFAX: (703) 816-4100

Does Not Comply  
Corrected Diskette Needed*See especially pp. 2, 23, 31*

ERRORED SEQUENCES FOLLOW:

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:36:23

INPUT SET: S26725.raw

45 (2) INFORMATION FOR SEQ ID NO: 1:

46

47 (i) SEQUENCE CHARACTERISTICS:

48 (A) LENGTH: 3018 base pairs

49 (B) TYPE: nucleic acid

50 (C) STRANDEDNESS: double

51 (D) TOPOLOGY: linear

52

53 (ii) MOLECULE TYPE: genomic DNA

54

55 (ix) FEATURE:

56 (A) NAME/KEY: misc\_feature

57 (D) OTHER INFORMATION: /label= FIGURE 8a

58

59 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

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62

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64

65 CTTACGAGAA CCTATCGATG ATGTTAAGTG AGGATTTTCT CTGCTCAGGT GCACTTTTTT

66

67 TTTTTTTTAA GACGGAGTCT CTTTCTGTCA CCTGGGCTGG AGTGCAGTGG CGTGATCTGG

68

69 GTTCACTACA ACCTCTCCCT CCTGGGTTCA AGCAATTCTT CTGTCTCAGC CTCCCAAGTA

70

71 GCTGGGATTA CAGGCACCCC CCGCCACACC CGGCTTATTT TTGTATTTTT AGTAGAGACA

72

73 GGGTTTCACT ATTGTTGTCC ATGCTGGTCT CGAAGTCGTG ACCTCATGTG ATCCACCCGC

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75 CTCGGCCTCC CAAAGTGCAG AGATTAGAGA CGTGATCCAC ATGGCCCAGC AGGACCACTT

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77 TTTAGCAGAT TCAGTCCCAG TGTTCATTTT GTGGATGGGG AGAGACAAGA GGTGCAAGGT

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79 CAAGTGTGCA GG TAGAGACA GGGATTTTCT CAAATGAGGA CTCTGCTGAG TAGCATTTTC

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81 CATGCAGACA TTTCCAATGA GCGCTGACCC AAGAACATTC TAAAAAGATA CCAAATCTAA

82

83 CATTGAATAA TGTTCTGATA TCCTAAAATT TTAGGACTAA AAATCATGTT CTCTAAAATT

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85 CACAGAATAT TTTTGTAGAA TTCAGTACCT CCCGTTCCACC CTAAGTAGCT TTTTGTCAAT

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89 AACATGTCAG CAGTTCTCAG CTTCTTTCCA GTGTTACCT TACTCAGATA CTCCCTTTTC

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91 ATTTTCTGGC AACACCAGCA CTTTCATGGCA ACAGAAATGT CCCTAGCCAG GTTCTCTCTC

92

93 TACCATGCAG TCTCTCTTGC TCTCATACTC ACAGTGTTTC TTCACATCTA TTTTGTAGTTT

94

95 TCCTGGCTCA AGCATCTTCA GGCCACTGAA ACACAACCT CACTCTCTTT CTCTCTCCCT

96

97 CTGGCATGCA TGCTGCTGGT AGGAGACCCC CAAGTCAACA TTGCTTCAGA AATCCTTTAG

global errorinsert  
cumulative  
base  
totals

etc.

Per 1.822(h) of Sequence Rules insert  
cumulative base totals at end of each line

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PATENT APPLICATION US/08/836,734DATE: 06/16/98  
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104  
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117 AGCTTCCTGC TTGCTGGCTG GGTTTCCCCC CCACGGAGGA GTCTCTCAC TCAGCACCTC  
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122  
123 TCAGCAAAAT CCAGAGGGAG AGCGCAGGAG GTGGGGTGAT TCTTATGCTC TGGCTCTTTC  
124  
125 TCTCTGAAAA AAAAAAAAAA ATCTTGCTTT TTATAAAAGT GGGTGGAACT CAGTTTAATT  
126  
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132  
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144  
145 AAGACCAAAA TTCCAGCTGT CTTGGAACCT AGGGTCTCTG AGGGAAGATG GGCATTGAAC  
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env*

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164 (2) INFORMATION FOR SEQ ID NO: 2:  
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## (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 11451 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: double

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: genomic DNA

## (ix) FEATURE:

(A) NAME/KEY: misc\_feature

(D) OTHER INFORMATION:/label= FIGURE 8b

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## (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

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185  
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189  
190 TGATAGTATT CCCTTCACAG GGCCAAATGG AATACTATCA GGAACACTAC ATAATGGAAC  
191  
192 TCAATAAATA ATAGCTACTG CGGCCGGGCG CGGTGGCTCA CATCTGTAAT CCCAGCACTT  
193  
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195  
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248 AGGAATTGGA CTCACATTGC AAAGGCACAG GGCAGGGCAG ATTTCTTACA GGTGTTAGGA  
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250 AGAACAACCC AGTTATGATC ACCTACTGCT CTGTCTCCAT TGAGGCCTAA AAAGGAAGTG  
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265  
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267  
268 CTTCCCAACC ATCTACCCGC AGCGGCAACA GTCGGCATGG ACCCCCTTAA GGCTTCAAGC  
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270 CTGGGAGGAA GCAGTTGCTT ATCTCTGGCT CCCTAATCCC TCCCCACCA CCTTCCACTA  
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272 TGTCCCAGAA AGACAGGAAG ACATCCTGTT TACTGTGGGT CTATTTTTGT CTTTGCAGCT  
273  
274 GTCTGGCTGC TTTTATTGCC TGCAGCCCTT CTCAAGTAGG TCCCTAAGAT ATTAGCACTG  
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276 TGACACCACA GGACCCTTCA GGTTGTACAG GAACCCCTGT CCAGGGCTCC TGTATACTTC  
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284 TGCTAGAGCA GAACCAAAC GAATTCTACC TGTGAGGGTC TCGTAGCTTC CGGGATGCTG  
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286 GGGAGTCAGC CTGTCTCCAG CTTCAAAGGC TCCCTCATGT CCCAGGATGA CCCACATTAT  
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404 CAAAGCAAAT TCTCTCCTCT TTTTCAATAT TTGTGGAAAA AGCAAGGTCT CCCTCTTGTA  
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424 CACCACACCC GGCCTAGTTT GATATTCTTA ATGTGCCCAA AGTATTCTCC TGTAACATTT  
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426 TTTAATAGCT ACACAATATT CAAACACACA GATATGTTAT AATTTATTTA CCAATACCC  
427  
428 TATTATTGGA AAGTTGAGTT CTTTTTTTTT TTTGTTTTGT TTTGTTTTGC TACTATTCTA  
429  
430 AAATGCTATA ACGAACATCC CAATAGATAC ATCTTTGTAT ACATCCATGG TGACTTCCAT  
431  
432 AGGACAGATT CCCAGCAGTA GAATTGCTGG GTTGAATGAT ATGCTTAGGG TAATGACAGA  
433  
434 AGAGTCATTT CAAGCAGCTT CCTAGGGTCT TAGAACTTAA GGATTAATGA GTCTTCCCGC  
435  
436 CCCCTCCCAG TCTATTGAGC ATGATCTGGA TCATGAGGAC TGAGATCTGG AAGAGACTGA  
437  
438 GATCTGGGAG AGGCTGAGAT ACCAAAAGCC CTGGCTCCAC CCATACCCCT CGCCCTGAAA  
439  
440 ACAGCTCTAG GAATTCCGCG GCCTAGCAAG GCTCCGGGAA GCTCCTTTTA AAGCTGTGAC  
441  
442 GTTAGTAGGC ACATGGACCA TAGAGACCTA TCCAGGGCTC ATGGGACTTT AGTGATCCTG  
443  
444 CCCTTCTCCC AAGGATCCCC CATGGCTGCA ACTTGGAAAT TTCTGCAAAT GGAAGAGCTA  
445  
446 CTCCTTAGGC ACGGTCATGT CTGAGCAGGG ATCTCCTCGG GCTTCTTAG AATTCTCTCC  
447  
448 CTGGGCACTG GGAATCTTGA TTTCTTGAAT ATTATGTTCC AGGTGGGTGT GGAGGAGGTG  
449  
450 AGGGGATGTA AAGAAGGCTA GACTTGGCCA GGCGCAGTGG CTCATGCCTG TAATCCCAGC  
451  
452 ACTTTGGGAG GCTGAGGCGG GTGGATCACC TGAGGTCAGG AGTTCGAGAC CAGCCTGGCT  
453  
454 AACATGGTGA AACCCCGTTT CTACTAAAAA TACAAAAAAT TAGCTGAGCA TGGTGGCAGC  
455  
456 TGCCTGTAAT CCCAGCTACT CGGGAGGCTG AGGCAGGAGT ATCGCTGGAA CACGGGAGGC  
457  
458 AGAGATTGCA GTGACCCGAG ATCGCGCCAC TGCACTCCAG CCTGGGCGAC ACAGCAAGAC  
459  
460 TCTGTCTCAA AAAACAAAAA AGAAAGAAAA AAAGGAAAAA CTAAGACTTA CATGTGTCAC  
461  
462 TTAACCCCTT TTCTCAAACC TCTTCTCTT CCAGGAATAG TCAACCCCTG GATGGCTTCA  
463  
464 GGGGAAGGGG GATCCTGAAG CCCAGGGCAG CCTCCAATC TACCCCTTCC TCCTTTGAAG  
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466 GATACTAAGG GGTCCAGAAA GGAGGGGCAG GAACTGTGTA CCCACCCAC ATCCCAGCAT  
467

*same*

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:36:36

INPUT SET: S26725.raw

468 CCACATTGCT CTCTGATGGT CAGGACAGAG CCTTCTCAGG GAGACCAGCC TGTCTGGAGC  
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470 TGTGTCTCTT GGCACCTCTTA AAGGGCCACT GAAGGTCCGT TCGTGGTCGT GAGGCACACT  
471  
472 TTCAGGGAGC AGAGTGGTCT GTGTCTTCAC AGAGCCCGGA AAATGAACTA GTATGAACTT  
473  
474 TGCCTCCAAG CAGCAGAACT TCTGTTCCCC CGCCCCTAAT GGGTTCTCTG GTTACTGCTC  
475  
476 TACAGACAAT CATTCCGGTT CAGTATGAGA CAAGAATGGC CTGCGGGCTG GTCAGAGGTC  
477  
478 ACGCCTACTC TGTCACGGGG CTGGATGAGG TAAGCCTGGT GGGGCTTGGT GGGGCAAGGG  
479  
480 CACCCTCCTG GGTAAACCTC ATGAAGTCAG GACTTAGCTG TTGGGGCCCC TGCCCTGTCT  
481  
482 GCAGAGCTTG CCTCCAATCA GGACATTCAG TTCAAGGTCC AAGCCACGCC TGGGAGCAGA  
483  
484 GGGGCCTGTG AAAGTGGTAG AGGTGGATCC TGCCACAGTT GGTGCACAGT TTATCTTTGC  
485  
486 TTTTCGTGCT AAAGATGGCA ATTTTCCAA CATTTCCAAT GAACAAATTG AAATATCACT  
487  
488 TAACTTTGCT TTTACAAAGT TGGTTTCATG TGTCTTGAG CTCCTGTTC TCTCGTGTC  
489  
490 AGATAGCTAC AGTTGTCTCT GGGTAGCCAC GGGGACTGGT TCCAGAAGCC CCAACAGTAA  
491  
492 CAAAATCTGC AGATGCTCAA GTCCCTTCTG TAAAATGGAG TAGTATTTGC ATATAACCTA  
493  
494 TGCACATCCT CCCATATACT TTAAGTCATC TCTGGATTAC TTACGATACC TAACACAATG  
495  
496 GAAATGCTAT GTAAATAGTT ATTGCACTGC ATTGGGTTTT TTTGGTATTA TTTTCTGTTG  
497  
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499  
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501  
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503  
504 CCCTGTACAG AGGGATGGGC TGAGAGGGGC AGTTGCCTGC ATCACCATT GCAGCAGACT  
505  
506 GGAGGGAGTC TGCTTGTTTG TAGTTCCTCA GTCAGCAGGG GCCTTTTGTC TTTCTTCCT  
507  
508 TTCTTTTTTT TTTTTTTTTG AGACGGAGTC TCACTCTGTT GCCCAGGCTG GAGTGTAGTG  
509  
510 GCACAGTCTC GGCTCACTGC AATGTCCGCC TCCTGGATTG AAGCGATTTT CCTGCCTCAG  
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512 CCTCCTGAGT AGCTGGGATT ACAGGCGCGT GTCACCATGC CCAGCTAATT TTTGTATTTT  
513  
514 TAGTAGAGAT GGGGGTTTCT CCATGTTGAT CAGGCTGGTC TCGAACTCCT GACCTCGTGA  
515  
516 TCCGCCCACC TCGGCCTCTC AAAGTGCTGG GATTACAGGC GTGAGCCACC ACGCCTGGCC  
517  
518 AGCAGGGGCC TTTTTTCTAA TTTATATGAA GACACCTAAT TTATATGTGT TAGCAAAGCC  
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520 CTCCTGTTTA TGCCTCACCT CCTCCCCGA AGCTCATACG GCAGGATGTT CCTGAGAAAA

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TIME: 13:36:37

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525  
526 CAGGCAGAAC ACCCTCGCGT AAGAGATTTG CCCCCAGCC CCGTCCCAGC CCTCAGCTAG  
527  
528 ACAGAAGATT CCCTTTCCAG AGAGGCTGCA GAGCATGAGA GCTCTTTCTG TGTGCTTAAG  
529  
530 GTCCCGTTCA AAGGTGAGAA AGTGAAGCTG GTGCGGCTGC GGAATCCGTG GGGCCAGGTG  
531  
532 GAGTGGAACG GTTCTTGAG TGATAGGTAG GTGAGGGGAC CCCACGGGAT TGGCGGTGGC  
533  
534 GGGGAACAGG GTCCGGGACA AGGCTGTGTT GGGAAGTGAG CCATGAGAGT ATTGAAGATG  
535  
536 CTTGGTATAA AATCACCCCTC AAAACCAATG ATCCGCAGAG AAGAGGGGCA CAGGTGTGG  
537  
538 CTCCAGGGAA GGGCCAGGAG TGGAAGCGGG GTGCTGGGGA CCCAGAGAGG TTGCTGACAA  
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541  
542 ATGAATGCAG GGTCTGCGC TAGAGAAGTG ACTTCCCTTC TTGGGGTCTT GTGTTGCCTT  
543  
544 TCCTGTGAAA TGGGAACAGT ATTATTAGCA CTTACCTTGT GGGCTGATAT TGAGGAGTAA  
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548 TCCCTTGAT TAAGGCACAA GGGCCCTTTG AAAAGAATTT TACCTGCTTT ATCAATTGAA  
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551  
552 CCATGCACAC TTCGTAAAAC CTCCATGGTC CTACTGGTTC CTGATTACCT CCACTCAATG  
553  
554 AGAGGCAATT CATTACTGAA TGAGCCATAA GCGCCTCTTA TTTGAGAGG GGGATGGCAG  
555  
556 GACTCAGTCG AGGAGAAGGA CCGCACCCAG GCAGCCTGGG CCCCTCGGCT CCTGTACTTA  
557  
558 TTTACTGCTG GGTACTTCCT AGCCCAGCAT GTAATTACTG GTTCGTTTCTG TCATTGCTTT  
559  
560 AGTAAATGTT TCTTGGGCAC CTACTACATA GGAGGCACAG GTCAAGGCAC TGGGGATATT  
561  
562 CTTTCTACCC ACCCCCTCCC TTGATACACT GTGATTAGGG ACTGACCGAT C  
563

*same*

564 (2) INFORMATION FOR SEQ ID NO: 3:

565

566 (i) SEQUENCE CHARACTERISTICS:

--&gt;

567 (A) LENGTH: 1834 base pairs

568 (B) TYPE: nucleic acid

569 (C) STRANDEDNESS: double

570 (D) TOPOLOGY: linear

571

572 (ii) MOLECULE TYPE: genomic DNA

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:36:39

INPUT SET: S26725.raw

573  
574 (ix) FEATURE:  
575 (A) NAME/KEY: misc\_feature  
576 (D) OTHER INFORMATION:/label= FIGURE 8c  
577  
578 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:  
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581 CGCGATCTTG GCTCACTGCA ACCTCCGCCT CCCGGGTTCA AGTGATTCTT CTGCCTTAGC  
582 CTCCTGAGTA GCTGAGACTA TAGGTGCCCC CCACCACGCC CAGCTAATTT TTGTATTTTT  
583  
584 ATTAGGACGG GGTTCACCA TATTGGCCAG GCTGGTCTCG AAATCCTGAC CTTGTGATCC  
585  
586 GCCCACCTCG GCCTCCCAA GTGCTGGGAT TACAGGTGTG AGCCATTGCG AGCAGCCCAG  
587  
588 AACTCAATTC TTAACCTTTA AAGTATGATG AGAAGAAGGA TCAAGCCCTC ACCAGCCCAT  
589  
590 TTAAGGAGTT TAGGCTCAGT CTTGAGGATG TGAGAAGTCA TTGCTATTGG GTTTCACACT  
591  
592 GAGGTAAACA GGTGAAGTCA GCATTTTGGT AGTTCACAGC AGCTGCAACT CTTTGTATTT  
593  
594 CTCTGATACC TCCTGTCCCA ACCTACATCA GGCCTTCCCT TCTTCCTGCT TCCTTAATTC  
595  
596 CTCCATTTTC CCACCAGATG GAAGGACTGG AGCTTTGTGG ACAAAGATGA GAAGGCCCGT  
597  
598 CTGCAGCACC AGGTCACTGA GGATGGAGAG TTCTGGTGAG TCCAGAACCC AGGAAGACCC  
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601  
602 TCAGATTCCA GCCCTTGGGA GATCTTGGCT GTGTTCTCCT CCAGCCCAAG GCCCAGCAAG  
603  
604 GATGAGGTTT TGAGAGGAGC CTTCCAGGCC ACAGGGACAA TGAGCCCAGG ACCAGGCCAA  
605  
606 CATGACATGG CTCTTGCCCT CTGTGTGCCC CTCCGCCACA CACTCTATTC CAGCCACAGG  
607  
608 CACCCTGGCC TTAGCACAAT TCTTTTCTGA GCCTAGGAAG CTCCACTTAC CCTGATCTTC  
609  
610 CAACGTCAAC CTCACCCTCT CTCAGGTTGT TTCTATTCAG GCTTCAAGTC TCAGCTTAAG  
611  
612 GAGAATTTTC AAGTCTCAGC TTAAGGAGAG CCCCCTAAGT TCCCCGAGGA CTGGGATTAA  
613  
614 TTTATGATGC TCATCACCTT TAAAATTGTT TGCTTAAGCC GGGCGCGGTG GCTCACGCCT  
615  
616 GTAATCCCAG CACTTTGGGA GGCCGAGGTG AACGGATCAC GAGGTCAGGA GATCGAGAAC  
617  
618 ATCTTGGCTA ACACGGTGAA ACCCTGTCTG TACTAAAAAT ACACAAAAAA AGTAGCCGGG  
619  
620 CGTGGCAGCG TGCGCCTGTA GTCCTAGCTG CTGGGGAGGC TGAGGCAGGA GAATCACTTG  
621  
622 AACCTGGGAG GCAGAGGTTA CAGTGAGCCC AGATTGCGCC ACTGCACTCC AGCCTGGGCG  
623  
624  
625

*same*

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TIME: 13:36:41

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628 GAAAGGTTTT GGGTGTTTTT ATTACTTTAT TTTTATTTA AAAACTATAA TAGAGACGGG  
629  
630 CCTCGCTATA TTTCTCGGGC TGGTCTCAA CTCTGGGCT CAAGCGGTCC TCCCACCTTG  
631  
632 GCCTCCCAAA ATGCTGGCAT GTGGGCCTGG TCAACATATG GGACCCCAAC TCTACAAAAA  
633  
634 ATTTTAAAAAT TAGCCAGATG TGGTGGCGTG TGCCTGTAGT CCCAGCTACT TGGGAGGCTG  
635  
636 AAGCAGGGGG TCACTTGAGC CCAGGAGGTT GAGGCTGCAG TGAACATGA TTGTCGTTCA  
637  
638 CTTTCTTCT GAACGTGAGA TTAAGTGTAG TCAGCAATTT GGCTTAGGAT TATTTATTCA  
639  
640 GAATTTTAA CCGTCACGTT GCGGCAAACC AGGT  
641

*same*  
↓

## (2) INFORMATION FOR SEQ ID NO: 4:

643

## (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 14664 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: double

(D) TOPOLOGY: linear

649

(ii) MOLECULE TYPE: genomic DNA

650

## (ix) FEATURE:

(A) NAME/KEY: misc\_feature

(D) OTHER INFORMATION: /label= FIGURE 8d

655

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

656

658 AGGAGGTGGA GGTTCAGTG AGCCAAGATC ATGCCACTGC ACTCTAGCCT GGGCAACAGA

659

660 GCGAGACTCT GTCTCAAAA ATACACACAC ACACACACAC ACACACACAC ACACACACAC

661

662 ACACACATAT ATATACACAC ATATATATAC ACACACATAT ACACACACAC ACGTCTGTAT

663

664 ATATATGTGT GTGTGTATAT ATACACACAC ACACTATTCT ATATATTCTT GTAGAGCTAT

665

666 GTGTGTCTCC TGTGCTATTG AGCATGAGCC CTTTTTTTTT TTTTTTTTTT TTGAGACAGA

667

668 GTCTCACTTT GTCGCCAGG CTGGCATAA ATGGCGCAAT ATCGGCTCAC TGCAACCTCC

669

670 GCCTCCTGGG TTCAAGTGAT TCTCCTGCCT CAGCCTCCCA AGTAACTAGG ATTACAAGTG

671

672 CCCGCCATAA TGCTCAGCTA ATTTTGTAT TTTCAGTAGA GATGGGGTTT CACCATGTTG

673

674 GCCAAGCTGG TCTCAAATC CTAGCCTCAG GTGATCCACC TGCCTCAGCC TCCCAAAGTG

675

676 CTGGGATTAC AGGCATGAGC CACAGCACCC TGGTGAGCAC TAGAGCTTAT TTCTTCTATC

677

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RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:36:42

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678 TAACTGTATT TTTGTATCCA TTAGCCACCC TCTTTTCATC CTCCCCTCTC CTTCCCTTCC  
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681  
682 CACATATGAG TGAGAGCATG CGACATTTAT CTTTCTGGCC CTGGCACATT TTTGAATCAT  
683  
684 TGTTAGAAAA GATGATGGTT TGGAGTAGAT ACATCAGAAG TGACAGCGTT TGCCCTAAAA  
685  
686 AGGAAAGACA GGCTCCTCTG GGACCCTGAC CAAGTTCCTG TGAACATTTT TATTATTGTG  
687  
688 CTGTGTTAGT CCTGGGGTCT TCCGTTCCCA GCCCTCCTCA CCTGCTCCCA TATGGCTCTC  
689  
690 TCTCTTCTTC CAACCTCTCA GGATGTCCTA TGAGGATTTT ATCTACCATT TCACAAAGTT  
691  
692 GGAGATCTGC AACCTCACGG CCGATGCTCT GCAGTCTGAC AAGCTTCAGA CCTGGACAGT  
693  
694 GTCTGTGAAC GAGGGCCGCT GGGTACGGGG TTGCTCTGCC GGAGGCTGCC GCAACTTCCC  
695  
696 AGGTGGGAGA TGCTCTTGAT GGGGGGAGGG TCTAAGCCGA AAAAGTTCCA GGCAGAAGAA  
697  
698 GCCTAACTAG TGCTTATTAA GTCTCTCTGT TCCAGACGTC CACTATCTTA TTAAACCTTC  
699  
700 CCTGTTTTAC TGAGAAGGAA ACCACCATGC TGAGAAGTTT GCAATAGGGA GCTGGGTAGC  
701  
702 AACTTTGGAA GCAGGAACCT GTGGGAACAA TGCAGATGCT GCTTGGACTT ACGATGAGGT  
703  
704 TATGTCCAGA TAAGCCCATC CATCTTTTGA AAATACCCTA AGTGAAAAGT GCATCCAATA  
705  
706 TGCCTAACCC CCCAAACCTC ATAGCTTACC CTGGCCTACC CTCAAACATT GCTCGGAACC  
707  
708 CTTGACCTTA AGCCTAAAGT TGGGCCAAAT CATCTAACTC CAAAGCCTAT TTTACAAAGA  
709  
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711  
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713  
714 GTAAAGTTAA AAAATTGTAG CCGAACCATC CTAAGTCAGG GACTGTGAGT ACTGTGTCAG  
715  
716 TAACAGTAAG GGCACATTTG GAGAACCAAG TTAGCAGCTG CTGCAATAGT TCAAGTCAGA  
717  
718 GATGATGAAA ACCTAGACCA AGTCAGTAGC AGCAGAGATG GAGGGGAGAC AGCAGATTTA  
719  
720 GGGAGAGCAT ATTGGGTGAT GTAGGGAAGG AAGAAGAATG ATGTCAAGAT TCCCAGTTGG  
721  
722 GGACCTGACA ACATTGCAAC ATAAGACACA CAAGAAGATC GGGTGGGTGG CTCATGCCTA  
723  
724 TAATCCCAGC ACTTTGGGAG GCAGAGCCAG GAGGATCACT TGAGCCCAGG AGTTCAAGAC  
725  
726 CAGCACAGGC AACATAGTGA CACCTCATCG TTACCCAAAA TAAAAAATAA AATGAGGTGG  
727  
728 GAGGATTGCT TGAGCTCGGG AGGTTGAGGC TACAATAAAC TGTGATCATG CCACTGCACT  
729  
730 CCTGCCTGGG TGACAGAGTG AGACCCTGCC TCAAAAAAAA AAGACACACA AGAGAAAAAT

*Same*

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TIME: 13:36:44

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734 CTTGGGACAT GGAAAGTTG AGGTTCTGT AGAGTGTCCT AGTGAAGATT TGTAATAGAG  
735  
736 CATCGGATGC GCATATTAGA TGGCACTTGG TGATATGATA AGAACTCAAA AAATATTTGA  
737  
738 GGAATAAAGG AAAGAAGAGG CCAGACGTGG TGGCTTATGC CTGTAATCCC AGCACTTTGG  
739  
740 GAGGCTGAGG CAGGCGGATC ACTTGTGGTC AGGAGTTCGA GACCAGCTTG GCTAACATGG  
741  
742 TGAAAACCCA TCTCTACTAA AGATACAAAA ATTAACCGGG GATGATGGTG GGTGCCTGTA  
743  
744 ATCCCAGCTA CTTGGGAGGC TCAGTCAGAA GAATCGCTTG AACCCAGGAG GCGGAGGCTG  
745  
746 CAGTGAGCCG AGATCGCGCC ACTGCACTCT AGCCTGGGCA ACAGAGCCAG ACTCCGTCTC  
747  
748 AAAAAAAAAA AAGTGAGAGA GATTGAGGCT GGGATATATG GCTCAGGCAT CATGCGCGTG  
749  
750 TAGGGGGCAG TTAAAAAGCA GAAGTAAGAA AGATTGCCCTA GGGAGGCAGG AAGGGTGAGG  
751  
752 TGAGAGGAGA AGAGGCCAG GACCAGATTC TAGTCACCAA CAGCGTTTAA GGGGCAGGTA  
753  
754 AGGAAAACAA AACCATCAGC AAAGACTGAG AATGAAAGCC CAGAGAGGAA GGAAAAGCCA  
755  
756 CACATACAAT CAGTACAGCT CCATCTGAAT AAAGGTAGCG CCCCCCCCCC CCCAAATCAT  
757  
758 TAGAGAAATG CCTGATTGCG TTTTCTGTGG ATTTTTCCTA AGAACCTAGA TGTGGGGAAT  
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760 AGAAATAAAT GGTTCCTCT GTCTCATCCC CTCCCTGCCC TCTGAGAGGA AGCTGTGATT  
761  
762 GCGTGCTCCC TTTCTGGGGG TGCAGATACT TTCTGGACCA ACCCTCAGTA CCGTCCGAAG  
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764 CTCCTGGAGG AGGACGATGA CCCTGATGAC TCGGAGGTGA TTTGCAGCTT CCTGGTGGCC  
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766 CTGATGCAGA AGAACCGGCG GAAGGACCGG AAGCTAGGGG CCAGTCTCTT CACCATTGCC  
767  
768 TTCGCCATCT ACGAGGTGTG TAGTCCTGAT TGGCTCCAGC CCAGGAAACA TACTTTCCCA  
769  
770 GAGAGGACGC TTCCAGGGGC TTCTAGAGGG GCCCTCTGCT TCCTCAATAC CAGTGACCCA  
771  
772 CAGAGCTCCT GGTATCAGGA CCACTTGTGT TTGTAACAAG CAAAAAATAC CAGGGGGGGC  
773  
774 ATTAGAGAGG CAGTGGAGCG GGCCTGGCAG AACAGGTGCC TGGGGGTCAG GCTTCCGCAT  
775  
776 GCGGGCTGCA GTTGCTGGCA TTGCCTTCCG CAGGCTCCTC ATCCTCATTC ACATCTGAAG  
777  
778 CATCTTCCTT TCTGTTTCTT CTCAAGGTTT CCAAAGAGGT ATAGCAGCAG CAGCGGCCAG  
779  
780 CAGTTGTGTG CAGCACTACC CAGGGGGGCC CGAGTCTGTC TGTGGCTCGT CGAGAAGCTT  
781  
782 CCTGGTGGGG TTTGTGGGCA GGACTTGTGA TAGGAGAGGG CCTTGCCTGT TGTTATTTCC  
783

*same*



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TIME: 13:36:45

INPUT SET: S26725.raw

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787  
788 TCCTCCACGC TTACAGCCAC ACACACAGTC ACACAGACGC GTTCTGAGGG TGGCTGCCCC  
789  
790 CTTGGGATGG AGGAATCACT TCCCTCAGAA CCCAGCCAAG TCCTCTAGGC CTCCTTGGGG  
791  
792 GTCCTTCCAG CCTGAGGGGC TTCGGAGCTG AGGACAGCTG TTCTGGTAAG TGTCCCTGAG  
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794 TGTGGGGATG ACACATTTC ATTCACTCTG AATCACAACA GAAAAGGGAA GAGGAATTGA  
795  
796 GGTAGGGAGC CTATTTAACC CTTGGGAGTC GGAAGTAGG GAGGTTGAAA CTGTGACATG  
797  
798 GGTGACCAGG GAGTTGGGAA GGGACCTTG GAGGTGGCTG TGGCAGGACA GGACGTTCTT  
799  
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804 ACATGCGGGA GGTGTCCCAG CGCTTCCGCC TGCCTCCCAG CGAGTACGTC ATCGTGCCCT  
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811  
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813  
814 GAGCACCTAC TATGTTCCAG GCACTGTCCT AGGCACTAAG GATAGAGTAG TGAAGTAAAC  
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816 AGAAAGAAAT CCCTGCCTTC ATGGAGCTTA ATATTCTAAC ATGAGACAAT AATGGATAGG  
817  
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820 AGAGGGATAG GAGGTGTTGG GGATGCTTGA AATTTTAGGT TAGCATGGCC AGGAAAGCCA  
821  
822 CATCCTGTCC CTGGCCACCA CAGATGAGCT CATAGCCCCT GCCACTCTGA TCTCTGTCCT  
823  
824 TGGAAGATGC ACCAGGTCCA TGGGTAGGTG GCTGGGTCAT GCCTTTGGGG GGCTCTGAGC  
825  
826 AATACTAACA AGAACCTGCG TGCCTGGGCT TGGCTGTCGG GGATGGTGCT GACATGGGGC  
827  
828 TGGTTCCTGG GGTGGGGTG TTCCAGGGGT TCTCTAGAGG CTGGTTCTGG CTTGGCTGCC  
829  
830 AGGAAGCCGT GCACCAGAGC AAACCGTCCA CGGGCCTCCT GCTTGCTTCT GGTGACACTG  
831  
832 AGACCCACCA TGTCTGTATT CCTCACAGGG AAGTTGAAAA TACCATCTCC GTGGATCGGC  
833  
834 CAGTGGTGAG TGGTTTAGAT CTTCTGTGCG AAAAGTCCAG AGGGTCCCCT TCCCTGACCA  
835  
836 TGCAGGGGAC AGATGGTGCA GGGGAGAATG GGCCTGGCA GAGGGAATGG GAGTCTGGGC

*same*

RAW SEQUENCE LISTING  
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TIME: 13:36:47

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841  
842 CTGGAAAGGA GAAGCAATTT GAACAATCGG AGGGAACAAG GCCACAGGAA GGGATGACAA  
843  
844 GAGCCGCAGC GAACACTGGA TTCTGAGACT GGATAACATT GGATTTTACA CATAGAGAAA  
845  
846 AGAAAGTAAG CTGGTGCCCG ACCTGGTGTT GACACTTGA TCCTCCACTT ACCAGCGGGG  
847  
848 TGACCTGGAC AATTTCTGTA ATCCCTCTCA CTCAGTTTCC TACTCAGTAA AACGGGGATG  
849  
850 ATAATGTGCC TTGCAAGGCT TTTGTGAGGC TTCATCAATG AGGTGATGTA TGTGAAGTGT  
851  
852 CTGGCACAGC ATGGGCACTC AAACAGAGGT GCTTTTTTAC ACTTTACACC TTACAAGGTA  
853  
854 CTTTTTACAT GTGTCATCGC GATACTTGCA AGGTTGCTGA GAGGTAGATG GGGTTATAAT  
855  
856 CCCTGGTGTT CAAGAAAGGA AGCAGAGGCT CAATGGGGTT GAATGACTTC TCTGAGTTCA  
857  
858 CAGAGCTCAG TAAGTGGCAG GGTGTTGGAAC TCACATTCAG ACTCTCTGAC TCCAGACTTA  
859  
860 GGTTTTTCCG CACCTCCACG CTGAGGCCAG CCCAGGCAG TGAGAAGCCC AAAGTCCGAA  
861  
862 GCACAGAGTG CTGTGTGTTG GGCTCTGTGT GTTGAGGAGT CTTGTGACTG CCTTGGGGCT  
863  
864 TTGGGCTGTA GTCAGCTGAC AGTCCTTTGT GCTCTGTGGG GATGACGTAG GCCAATGGGA  
865  
866 GGACAAATGC CCCTCTGAAC TGTCTTCTGG GCAGTGACAG TCATGGTCAT AATCCTGACC  
867  
868 CTGAGCCAGT GCCAGGTCTC CAAGTGCCTT CTGAATGACC ACAGGCGATT GGTTTTAGTG  
869  
870 GTAGGTGCGT GGGGATCTGT TCTGGTCATC TGGATGCTGG TCATCGGGTG CAGTATTGAT  
871  
872 CAGGACCTGC AAACCCAAAA GCTTATGGGA GCTGGCACGT CACGTGAGTA GAGCAGGCAG  
873  
874 GTGCAGGGTT TTTGATGTCC CTGCACTGAC ACAGTTGTCT GCAGTTCTCC AATTTGACAT  
875  
876 TTGGGCTCCA GTGTCGAGGG TCAAACAAGG AATTTTGGGG CGTGGGCCAA ATCTGGGAAG  
877  
878 ACACAGGGAG CAGGGCCCTT TGGCTCAAGC TGATAGTTGC CGCAGGGATT ACCAGGCCCA  
879  
880 GGGCAGCCTG CCACAAGCTG GGGCTTTTAC CAAAGAAAAT CTCCCTATGT TAAATGCTTG  
881  
882 CTCAAAAATT TTTAAAAAAT ATTCTGTAAG TCAAAATCCA TTGTTAGGTC AGTTTGAGAG  
883  
884 AGCCATGTTT TTGGTGTTTT AGTAACCAAT TTCATTTTTT TATTATTTAT TTATTTGTTT  
885  
886 ATTTTTGAGA CGGAGTTTCA CTCTTGTCAC CCAGGCTGGA GTGCAATGGC ATGATCTCAG  
887  
888 CTCCTGCAA CCTCCGCCTC CCGGGTTCAA GCAATTCTCC TGCCTCAGCC TCCTGAGTAG  
889

*same*

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:36:48

INPUT SET: S26725.raw

890 CTGAGATTAC AGGTGCCCCAC CATCACGCCT GGATAATTTT TGTATTTTTT AGTCGAGATG  
891  
892 GGGTTTCACC ATGTTGGCCA GGATAGTCCT GAACTACTGA CCTCAGATAA TCCGCCCACC  
893  
894 TCAGCCTCCC AAAGTGCTGG GATTACAGGC ATGAGCCAGC ACGCCCGGCC ACCAATTTCA  
895  
896 TTTTTTAAAA AAGGAAGAAA GAAAACCTTA GCCAGAAGAT CTTTTTCCTT GCCATATGCA  
897  
898 GTAAGAGTAG ATTATAAAAA CAAAGTCAGA GCAGTCACTG GTGTCTGGGC ATGGAGGAGA  
899  
900 AAGAAGAATT CTCTTCTCCC TTCACCTCC ATGCCCCTTT TTGGCTCCAT GTGATTCAGA  
901  
902 TTTCTGGACC CTGGAGCCCC ACCCCAAGCT AAAGACCAGG ATACAGGGAA GCCACAACCA  
903  
904 CTGGCGGTTC TGAGAACTTA CTTTTCACTT ATTCTGCATT TACTGTTTCC TTTTCTTATG  
905  
906 CAGAAAAAGA AAAAAACCAA GGTAGGTGTG TGGGTAGAGA GCATGAAGTG TGTGTACTCA  
907  
908 TGCATATGTA TGTGCATGCA TGTGAAGTGT GCATGTGTGA GCTCATATGC ATCCATGCAC  
909  
910 CAGACTTGCC TCTTCTCCC CCTCCTTCTT GAGCTTCTGC TGGGGCCGAG CGTGCAGTAA  
911  
912 TGACAACTAC GATTTGCTGG GGGAAGGCTA CGTGCCAAGC ACTCTTTTAG GTGCTTTCCA  
913  
914 TGATTAATTC CTTCTCACA ACAGCCCTAT GAGATTAGTA CTATAACTAT CCCCATTTTC  
915  
916 AGAGGGAGAA AAGGTACAGA CTTGACTAAC TTGCCCAAGG CCACACAGCC AGAGAGGGGC  
917  
918 AGAGCCAGTA CTTAGAGCCA GGCAGTCTGG GTCCAGAGTC CGTGCCTGA ACCACAAGAG  
919  
920 GCCATCATAC GCCATCAGAT TTGGTGCTAG CATTTCTGGT GGTGCCTGGT GGTGATGGAT  
921  
922 CCATCACAGG GGTCTCCAG GTCAGTGTGC TGGCCCAGAC CAGAGCTGAC ACTCTCAGG  
923  
924 CACTACCACA TTCCAGGCAC TGTGCTTGGG GTCAGTCCCT CTCTTTTTTT TCCCCCCAA  
925  
926 TTATAACAGT ATCTACAAAG TAGGTGCTGT TATTTTCCC CTTTCACAGG TGAGATAGAC  
927  
928 TCAAAGAAGT GAACTTGCCC AAGGAACAGA ACTAATGAGT GGGGAAAATG GAACTGGAAA  
929  
930 CCATGTCTGT TTA CTCCAAA ACCTGTGTTT CTTGCCCTCT TTCTCTGATG CCAGCCCCCT  
931  
932 ACACTTCAAG GCCTGTGTTG TCCAGACCCA CACTCGGGCC TGCCAGTGTG TGCCTGGCAG  
933  
934 GGATGCTCCA TGGCCACACC ATATCCATCC TACACATCCC CCCTCAGACT GTGACCTCCA  
935  
936 TTTGCTCTGG GATCCCCACA AGCTTCAGCT GCTTGAGCAA GACACTGCTT AGAAGGCAGA  
937  
938 GCAAGCCAAG GCCTCTGGGG CCTGCTGGGA GCCAAAGCTG GGGAGCCGTT TCCACGGGTC  
939  
940 TATCTGCTTG AGCTGTCCTA GATGAGCAGC ATGGAAGGGC AGTGGTGCAT GAGTCCAGGC  
941  
942 GGGCTGCTTT TCTGCTCCGA GAGGCTCTGC CTGCCAGTT GTTCTCTGCA TTGCAGCCTC

*same*

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:36:50

INPUT SET: S26725.raw

943  
944 AATCCCCACA GCCTTGCCTT CCCCCGGCTT TCCCTACAGG TGCACCGCAT CCACAGTGTT  
945  
946 GGCACCATGC AGCAGCCGCT CTCGTCCTT TTCATATCCT TGTCACCTGC ACGAGCATGT  
947  
948 CTTGAAAATA TCCCTTGTTT GTGTAGCATC TTAAATGTTT TTGCAGTATG ATTTTGCATT  
949  
950 CAGTATCTCA TTTGATCCCC ACAAGAGCCC TATGAGGAGG GAAAGCAGAT TTTACCATTA  
951  
952 AAGGATGAGT AAAGTGGGC CAGAGAGGAT ATTTTGGTT TTTTGTGAGA CAGTCTCACT  
953  
954 CTGTACCCCA GCCTGGAGTG CAGTGGCTTG ATCTTGGCTC ACTGCAAGCT CCACCTCCCA  
955  
956 TGTTACACC ATTTTCCTGC CTCAGCCTCC CAAGTAGCTG GGACTACAGG CACCCACCAC  
957  
958 CACCCCAGC TAATTTTTTT GTATCTTTAG TAGAGATGGG GTTTCACCCA GTTAGCCAGG  
959  
960 ATGGTCTTGA TCTCCTGACC TTGTGATCTG CCTGCTTCGG CCTCCTAAAG TGCTGGGATT  
961  
962 ACAGGCGTGA ACCCCCCCTGC CCGGCCAGAG AGGATATTTT TTAATGAGGG GCAGGGCTGG  
963  
964 GATTCCAGCC CAGTGTTCTG ATGGCTCACC CACTGACCAT TCCACTAATC CGTGTCTTTT  
965  
966 TTCAATCTAA ACTTTCAGGG TTGTAGAGGT TCCTTTGAGG TGCCTCAGTA CTTCCATGGT  
967  
968 GATGTGGGGT CTGAGGGCCA AGAGCTCTGT TCTCATTAAT CAGAGAAGCT TGTGTTTTTA  
969  
970 AAAACACCAT GTTTACTGCA GGAAATTTAA TTGGACAGTG TTTCCATCTG GAAAAAAAAA  
971  
972 AGTCTACAAA ATACTTGACA ATCACTGCAC TAGATCATGC TGCTTTTAGC ATTCTTAGCA  
973  
974 TTTACGTGC TGAGCTCTCA ATACTCTACC ATGAGGAGGG ATGGAGTGGG TATGAAAAGA  
975  
976 TAAAGAACTG AAGTCACACG GCTTGTCTAGT GGCAGAGATA GAGCTTGAAC CGAGGTTGAA  
977  
978 GAGCTCCCGC CTATTCCTTT CCTCTTCTCA CTGGATAAAG CTGCTCCAAG AGAGGTGCTG  
979  
980 CCTCAGTGTG CCTGTTCAGA CTGTAATCCT CCCTTCCTTC CTGCCTCCTC CCTCCTCTCT  
981  
982 CCAGCCCATC ATCTTCGTTT CGGACAGAGC AAACAGCAAC AAGGAGCTGG GTGTGGACCA  
983  
984 GGAGTCAGAG GAGGGCAAAG GCAAAACAAG CCCTGATAAG CAAAAGCAGT CCCCACAGGT  
985  
986 GTCTGGGCAT GTGGCATGGG TGGGGTGGCC AGCACGCTAC AGGGGCTTCC TATGCGCTTG  
987  
988 GGATACACAG GGGCTGGAGG CTCCCAGGA GTTTGTCTTG AACATCTGGA GGTTTGAATT  
989  
990 TGTCCCCTG ACCTTTTCTT TCAGCAAGTT CCCCTGAAAT TTGGGCTGCT GCTTGGGTGA  
991  
992 ATATCCCAGG ATGGGGGTTC CATTCTAGGA GTGGACTGGC AGGCTGAGCC TCCCATGGAG  
993  
994 CTGATCCAGC CAGGATACAG AGAAGGGGAG GCAAAGGCTG AGACAGAACC AGCTTGAGAG  
995

*same*

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:36:52

INPUT SET: S26725.raw

996 CGGAGGCGCA ACTCTTGTCT CCTGGTGGCC TTGAGCATTT CACAATAGGG GGATAAAGGA  
997  
998 TAGGAGCAGA AAAGTGGGGC TGACTTCAGA AATGGGGTCC TCTAGAGCTC ACGGGAGGGT  
999  
1000 GTTAGATTGG AGTGGGAGCT TAGTGGAGGT GAGCCTTAGA GGCAAAAGTC TCCAGACCAA  
1001  
1002 TCCAGGCCCC CTCTTCTATC CGGGGGCCCC TCTTCTATCC AGGGCCCCCTC TTCTGTCTGG  
1003  
1004 GAGCCCCCTCT TCTATCTGGG GCCTCATGCA GTGGGGCCTA GGGGAGGTTC TCTGAGGACT  
1005  
1006 TGGCCTTGAT GACAGGGTGG CTGGAGGAAT CAGAACGGTC AGACCTTCTT TGACCTGCGG  
1007  
1008 GCACCTTTAG TTGGAATGCT CAGGCCTGGG ATGGTGGAGG GGGCTCTTGC AGGTGGGGAC  
1009  
1010 TGGGGTGGCG GGGAGGAGGC TGTATGGCCG CCATATCTCC TTTGGCTGGG GGCCTCAGGG  
1011  
1012 CTGGAGAGGT GTGAAGAGTC CCTGAGGCCT CGATGCATCT CACTCCAGCT CACCAGGTCT  
1013  
1014 GCATTTGCCC GTCCCCAGCT CCTGCTGCCA CCCCCGGCCG TTTTAGGCAC TTGGCTCCCT  
1015  
1016 TGGCCCAGAG GAGCTTGCCCT CACAGGCCTG TGCACCTCTG ACCCCTGTGA ACCAGTTTTC  
1017  
1018 CTTTGTGCCT CCACAGCCAC AGCCTGGCAA CTCTGATCAG GAAAGTGAGG AACAGCAACA  
1019  
1020 ATTCCGGAAC ATTTTCAAGC AGATAGCAGG AGATGTGAGT ACCTCCAAGC CCAGGACGCC  
1021  
1022 CACAGGTGCT TCCTTCTCTC CTGGATTAAC TGCTCAGATT ACCAATTATT TCATTATTGT  
1023  
1024 TTGGTAGAGG TCACTTTGGA CTTGCGGTGA GCCAGGGGAT GTGTGCGTAG CACACAAATC  
1025  
1026 CACAAGCCCT TGAGTTTTGG ACTGCCACGT CTGCTGGGGG GCTCAGAGGC CTTTTTGCTC  
1027  
1028 TGAGCTGCCC ACGGTGGTCC TGATAGCTGA GGTGCAGTAT CTGGCCCCCT GTCTTCCTCA  
1029  
1030 GAAAAGCCCC AGCTTCCCAT GACATAATAG CACCGACAGG GATTTTACAA ACACAGCCAG  
1031  
1032 GTGGAATTTG TTTTGCAAAG TGTCCGCGCC AGGAGCTGCT GTACTCCTGA ACCATGACCC  
1033  
1034 TCCTCTCCCT TCCTCCTCAG GACATGGAGA TCTGTGCAGA TGAGCTCAAG AAGGTCTTAA  
1035  
1036 ACACAGTCGT GAACAAACGT GAGTTGCTCA AACCAAATGG GGGTGGGGTG GGTGGGGAGT  
1037  
1038 CCCGTTGTCT CAAAGCAGCT CCTCACTCTT CTCCATCCCC CCAGACAAGG ACCTGAAGAC  
1039  
1040 ACACGGGTTT ACCTGGAGT CCTGCCGTAG CATGATTGCG CTCATGGATG TATCCTTCCT  
1041  
1042 GCGGCCCCCT CCCGACCCTC TGTCATCAGC CCACGGGGGC CAAGGCAACA TACAGGGTGC  
1043  
1044 CCAGTCAGGC AAAGGGCCCT AATTTGTGCC CAGGGAACT TAAGGAGACC CTGATTGAGA  
1045  
1046 ACATCTTGGA TACTCGTCTG AAAGGGGTTG TTAGAGGCGG AAGGGGAGGA TGTGGGGTTG  
1047  
1048 TAACTGCCCT AACCCTGTG CTTCTCTCAG GCCTGGGATC CTGCCCAAGC AAAAGTGGTC

*same*

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:36:53

INPUT SET: S26725.raw

1049  
1050 CTTAGGAGAG CGGCTCCTGG GTTACAGAGT AGGCGCAATC TCTGACTGGT GGTGGAGTGG  
1051  
1052 AGGGGAGGGT TAAATAGTAC AACAGGGCAG TGGGTAGGAC AGCCCGGAGT CTCCTAGACC  
1053  
1054 CTCCCTCCAA ATCCAGGGGG ATTTTGCTGT GTGCTGTGTA GCCCTGACCT CCCTCCTCCA  
1055  
1056 GACAGATGGC TCTGGAAAGC TCAACCTGCA GGAGTTCCAC CACCTCTGGA ACAAGATTAA  
1057  
1058 GGCCTGGCAG GTGGGAAGAG AAAATGAAGC GTGGGAGTCA AGAATGGGGT TGATTTGGAG  
1059  
1060 ATTCAAGTGTG TGACCTCCAT CCTCAAATTT TCTATTGCCA GAAAATTTTC AAACACTATG  
1061  
1062 ACACAGACCA GTCCGGCACC ATCAACAGCT ACGAGATGCG AAATGCAGTC AACGACGCAG  
1063  
1064 GTGCTGAGAA GGAAGGGGTG TCAGGGATGT GGACCCGAGA CGGTGGGAGC AGGAATGGGA  
1065  
1066 GGGGACTAGC TACTAGGGCC CCACTAGAGA AGGAGAGGGA AAGGGCTTCT CACTTTCCCT  
1067  
1068 TCCCAGGTCA CAGAGTGTCC GAGAGGCAGG GAAAATAGAA GACAGGCCCA AGGCCTCCAG  
1069  
1070 CTCCACGTCC ACCTCTAACA TGGTCCCCTC CACAGGATTC CACCTCAACA ACCAGCTCTA  
1071  
1072 TGACATCATT ACCATGCGGT ACGCAGACAA ACACATGAAC ATCGACTTTG ACAGTTTCAT  
1073  
1074 CTGCTGCTTC GTTAGGCTGG AGGGCATGTT CAGTAAGTGG GAGAGGGGGG CTGCCCTCTG  
1075  
1076 CTCTCTTGCA GGGGCAGTTG TGGCAACAGG CATCTCACCT GATAATCTCC AGTCTGCTCC  
1077  
1078 ATCCAGGCTG AACAAGGGCC AATGACCTCT TTAGGCCAG AATGGGATGG CAAAGGGAGG  
1079  
1080 GTTACTGGTG ATTCTCTGCC TGCACATCTT TGTGCTGATG AGGGACAGCA CTGGGCACAC  
1081  
1082 GGTCTCTGA GGGGAAGTTA CAGTAGTAGA GGCGGAGTGC GCCTGTAACCT GGCCTCTGGC  
1083  
1084 CTGTGCATTC TTTCACAGGA GCTTCTCATG CATTTGACAA GGATGGAGAT GGTATCATCA  
1085  
1086 AGCTCAACGT TCTGGAGGTA AAGCATAGGC ACAGCACATT CCCCCTACAC ATTAAAACTC  
1087  
1088 AAGGTGGAGG GGTCAACGGG GCGGACTGGA CCCAGGGTGT GCTCCTCATT TCCACACAGT  
1089  
1090 GGTGGAGGGA AGGGATAGGA ACAGAACATG GAGGGAGGCT CAGCAGGCTC CCAGGACACA  
1091  
1092 TGCACTTGAG GCCCAAAGG ACCTCTGCTC CCCAGTCAC TTGATGCGGG AAAACATGCA  
1093  
1094 CCTTCTTAGG GAAGATCTAG GAGAAAGGAA ACAGTAAGCC ACTGCTTCTT GGAAAATCTT  
1095  
1096 CTGGGGGTCT GACCTGCTGG GACTGTTCCC TTTCTCTTG CCCCCTAAGA TTCTTAGGGC  
1097  
1098 GGGGGGGGGG GGGGGTCACT CTTTTCTGAT CTACATTCTG ATCTTGGGAC TTCTTTAGT  
1099  
1100 GGCTGCAGCT CACCATGTAT GCCTGAACCA GGCTGGCCTC ATCCAAAGCC ATGCAGGATC  
1101

*same*

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:36:55

INPUT SET: S26725.raw

1102 ACTCAGGATT TCAGTTTCAC CCTCTATTTT CAAAGCCATT TACCTCAAAG GACCCAGCAG  
1103  
1104 CTACACCCCT ACAGGCTTCC AGGCACCTCA TCAGTCATGT TCCTCCTCCA TTTTACCCCC  
1105  
1106 TACCCATCCT TGATCGGTCA TGCCTAGCCT GACCCTTTAG TAAAGCAATG AGGTAGGAAG  
1107  
1108 AACAAACCCT TGTCCCTTTG CCATGTGGAG GAAAGTGCCT GCCTCTGGTC CGAGCCGCCT  
1109  
1110 CGGTTCTGAA GCGAGTGCTC CTGCTTACCT TGCTCTAGGC TGTCTGCAGA AGCACCTGCC  
1111  
1112 GGTGGCACTC AGCACCTCCT TGTGCTAGAG CCCTCCATCA CCTTCACGCT GTCCCACCAT  
1113  
1114 GGGCCAGGAA CCAAACCAGC ACTGGGTTCT ACTGCTGTGG GGTAAACTAA CTCAGTGGAA  
1115  
1116 TAGGGCTGGT TACTTTGGGC TGTCCAACCTC ATAAGTTTGG CTGCATTTTG AAAAAAGCTG  
1117  
1118 ATCTAAATAA AGGCATGTGT ATGGCTGGTC CCCTTGTTT TTGTTGTCTC ACATTTAGAT  
1119  
1120 ATCAGCCATG CATGACTGAA TGGCTTCCAA TCATATACTC ACCTATCACC TACAAGAGAA  
1121  
1122 CAATGAAAAA CACACACAAA AACAAAATCT TGAATTTTGT AATCATGCCT ATTGCTATTT  
1123  
1124 CTTGAGCATA AGAATGGCTC AGATACTTTC CAAGACATAA AAGGAAGGCA GAGGAATAGT  
1125  
1126 TGTGCTGTA AAAGACATCA AGAATAAATG GGGTCATGTA CAACGGGAGG GGCCGGTTAC  
1127  
1128 CTGAATAATG GAGTGGAGAT TGAGCTATCC TAGCTCCTCT GCTCACTAAC TGACCTGTCTG  
1129  
1130 CATGACCGTG GACAAAACCC TGAACGCAGC TGTTTGTTTG CTAAACTTCT CTGGACCATG  
1131  
1132 GCCTGCGGCA TATCTATAGG CATCCTGTGT TTTCCACCCA GTTTCCTTCT TCCTCGCTAA  
1133  
1134 GCCAACGTGG AAAGGGCTGG CCGTGAATAT GCAGACAAGG TAACGAAAGT AAACCGTCAA  
1135  
1136 TTAGTAAAG TACTTCATTT TCCTCTTGTA TTTGCTTCAT TCTTGCTTCA CAAAGTTACG  
1137  
1138 AAGTCCACAG CTTTATACCA AAATGTAAGA AGGCTATTTG CTTATAAACA TTTTGAGTCA  
1139  
1140 GGTGTCATCT GATTTTATTC TTCTAATCCA TATTCAATAT TAAAAAATCA GAAACCAAGG  
1141  
1142 GTGCTGGAGC AGCTCTAGGG CATATATTTT TCTTAAATAG GAGAAAGATT TTCAACAGCT  
1143  
1144 TTTCTCCTT GACCCCTCC TTTCCCAATT TATTTGGGTC ACTACCTTGA ATTTAGAGTG  
1145  
1146 AATCTGGGAA ATGTAGTCAC CAGG  
1147  
1148

*same*1149 (2) INFORMATION FOR SEQ ID NO: 5:  
1150

1151 (i) SEQUENCE CHARACTERISTICS:

--> 1152 (A) LENGTH: 5149 base pairs  
1153 (B) TYPE: nucleic acid

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:36:56

INPUT SET: S26725.raw

1154 (C) STRANDEDNESS: double  
1155 (D) TOPOLOGY: linear  
1156  
1157 (ii) MOLECULE TYPE: genomic DNA  
1158  
1159 (ix) FEATURE:  
1160 (A) NAME/KEY: misc\_feature  
1161 (D) OTHER INFORMATION:/label= FIGURE 2  
1162  
1163 (ix) FEATURE:  
1164 (A) NAME/KEY: misc\_feature  
--> 1165 (B) POSITION : 1303..3764  
1166 (D) OTHER INFORMATION:/note=CDS  
1167  
1168 (ix) FEATURE:  
1169 (A) NAME/KEY: misc\_feature  
--> 1170 (B) POSITION : 1631  
1171 (D) OTHER INFORMATION:/note= CGA->TGA ; nCL1 mutation in one  
1172 LGMD2A family  
1173  
1174 (ix) FEATURE:  
1175 (A) NAME/KEY: misc\_feature  
--> 1176 (B) POSITION : 1848  
1177 (D) OTHER INFORMATION:/note= CTG->CAG ; nCL1 mutation in one  
1178 LGMD2A family  
1179  
1180 (ix) FEATURE:  
1181 (A) NAME/KEY: misc\_feature  
--> 1182 (B) POSITION : 1853  
1183 (D) OTHER INFORMATION:/note= CAA->CA ; nCL1 mutation in one  
1184 LGMD2A family  
1185  
1186 (ix) FEATURE:  
1187 (A) NAME/KEY: misc\_feature  
--> 1188 (B) POSITION : 2004  
1189 (D) OTHER INFORMATION:/note= GGG->GAG ; nCL1 mutation in one  
1190 LGMD2A family  
1191  
1192 (ix) FEATURE:  
1193 (A) NAME/KEY: misc\_feature  
--> 1194 (B) POSITION : 2248  
1195 (D) OTHER INFORMATION:/note= CGG->CG ; nCL1 mutation in one  
1196 LGMD2A family  
1197  
1198 (ix) FEATURE:  
1199 (A) NAME/KEY: misc\_feature  
--> 1200 (B) POSITION : 2364  
1201 (D) OTHER INFORMATION:/note= GTG->GGG ; nCL1 mutation in one  
1202 LGMD2A family  
1203  
1204 (ix) FEATURE:  
1205 (A) NAME/KEY: misc\_feature  
--> 1206 (B) POSITION : 2382

*G/o badly change this  
to "LOCATION:"*



RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:36:58

INPUT SET: S26725.raw

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1207          (D) OTHER INFORMATION:/note= TGG->TAG ; nCL1 mutation in one
1208 LGMD2A family
1209
1210      (ix) FEATURE:
1211          (A) NAME/KEY: misc_feature
--> 1212          (B) POSITION : 2771
1213          (D) OTHER INFORMATION:/note= CGG->TGG ; nCL1 mutation in one
1214 LGMD2A family
1215
1216      (ix) FEATURE:
1217          (A) NAME/KEY: misc_feature
--> 1218          (B) POSITION : 3018
1219          (D) OTHER INFORMATION:/note= GGG->CAG ; nCL1 mutation in one
1220 LGMD2A family
1221
1222      (ix) FEATURE:
1223          (A) NAME/KEY: misc_feature
--> 1224          (B) POSITION : 3372..3373
1225          (D) OTHER INFORMATION:/note= Deletion AC ; nCL1 mutation in one
1226 LGMD2A family
1227
1228      (ix) FEATURE:
1229          (A) NAME/KEY: misc_feature
--> 1230          (B) POSITION : 3533
1231          (D) OTHER INFORMATION:/note= AGC->GGC ; nCL1 mutation in one
1232 LGMD2A family
1233
1234      (ix) FEATURE:
1235          (A) NAME/KEY: misc_feature
--> 1236          (B) POSITION : 3609
1237          (D) OTHER INFORMATION:/note= CGG->CAG ; nCL1 mutation in one
1238 LGMD2A family
1239
1240      (ix) FEATURE:
1241          (A) NAME/KEY: misc_feature
--> 1242          (B) POSITION : 3616..3619
1243          (D) OTHER INFORMATION:/note= Deletion AGAC ; nCL1 mutation in one
1244 LGMD2A family
1245
1246      (ix) FEATURE:
1247          (A) NAME/KEY: misc_feature
--> 1248          (B) POSITION : 3665..3666
1249          (D) OTHER INFORMATION:/note= AG->TCATCT ; nCL1 mutation in one
1250 LGMD2A family
1251
1252
1253
1254
1255      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
1256
1257 ATATCAGTTA GCCTGGTTTC ACTATACAGT ACATCATTTT GCTTAAAGTC ACAGCTTACG
1258
1259 AGAACCTATC GATGATGTTA AGTGAGGATT TTCTCTGCTC AGGTGCACTT TTTTTTTTTT
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RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:00

INPUT SET: S26725.raw

1260  
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1262  
1263 TACAACCTCT CCCTCCTGGG TTCAAGCAAT TCTTCTGTCT CAGCCTCCCA AGTAGCTGGG  
1264  
1265 ATTACAGGCA CCCCCGCCA CACCCGGCTT ATTTTGTAT TTTTAGTAGA GACAGGGTTT  
1266  
1267 CACTATTGTT GTCCATGCTG GTCTCGAACT CGTGACCTCA TGTGATCCAC CCGCCTCGGC  
1268  
1269 CTCCCAAAGT GCAGAGATTA GAGACGTGAT CCACATGGCC CAGCAGGACC ACTTTTTAGC  
1270  
1271 AGATTCAGTC CCAGTGTTCA TTTTGTGGAT GGGGAGAGAC AAGAGGTGGC AAGGTCAAGT  
1272  
1273 GTGCAGGTAG AGACAGGGAT TTTCTCAAAT GAGGACTCTG CTGAGTAGCA TTTTCCATGC  
1274  
1275 AGACATTTCC AATGAGCGCT GACCCAAGAA CATTCTAAAA AAGATACCAA ATCTAACATT  
1276  
1277 GAATAATGTT CTGATATCCT AAAATTTTAG GACTAAAAAT CATGTTCTCT AAAATTCACA  
1278  
1279 GAATATTTTT GTAGAATTCA GTACCTCCCG TTCACCCTAA CTAGCTTTTT TGCAATATTG  
1280  
1281 TTTTCCATTC ATTTGATGGC CAGTAGTTGG GTGGTCTGTA TAACTGCCTA CTCAATAACA  
1282  
1283 TGTCAGCAGT TCTCAGCTTC TTTCCAGTGT TCACCTTACT CAGATACTCC CTTTTCATTT  
1284  
1285 TCTGGCAACA CCAGCACTTC ATGGCAACAG AAATGTCCCT AGCCAGGTTC TCTCTCTACC  
1286  
1287 ATGCAGTCTC TCTTGCTCTC ATACTCACAG TGTTTCTTCA CATCTATTTT TAGTTTTCTC  
1288  
1289 GGCTCAAGCA TCTTCAGGCC ACTGAAACAC AACCCCTCACT CTCTTTCTCT CTCCCTCTGG  
1290  
1291 CATGCATGCT GCTGGTAGGA GACCCCCAAG TCAACATTGC TTCAGAAATC CTTTAGCACT  
1292  
1293 CATTTCTCAG GAGAACTTAT GGCTTCAGAA TCACAGCTCG GTTTTAAAGA TGGACATAAC  
1294  
1295 CTGTCCGACC TTCTGATGGG CTTTCAACTT TGAACGGAT GTGGACACTT TTCTCTCAGA  
1296  
1297 TGACAGAATT ACTCCAATT CCCCTTTGCA GTTGCTTCCT TTCCTTGAAG GTAGCTGTAT  
1298  
1299 CTTATTTTCT TAAAAAGCT TTTTCTTCCA AAGCCACTTG CCATGCCGAC CGTCATTAGC  
1300  
1301 GCATCTGTGG CTCCAAGGAC AGCGGCTGAG CCCCAGTCCC CAGGGCCAGT TCCTCACCCG  
1302  
1303 GCCCAGAGCA AGGCCACTGA GGCTGGGGGT GGAAACCCAA GTGGCATCTA TTCAGCCATC  
1304  
1305 ATCAGCCGCA ATTTTCCTAT TATCGGAGTG AAAGAGAAGA CATTCGAGCA ACTTCACAAG  
1306  
1307 AAATGTCTAG AAAAGAAAGT TCTTTATGTG GACCCTGAGT TCCCACCGGA TGAGACCTCT  
1308  
1309 CTCTTTTATA GCCAGAAGTT CCCCATCCAG TTCGTCTGGA AGAGACCTCC GGAAATTTGC  
1310  
1311 GAGAATCCCC GATTTATCAT TGATGGAGCC AACAGAACTG ACATCTGTCA AGGAGAGCTA  
1312

*same*

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:01

INPUT SET: S26725.raw

1313 GGGGACTGCT GGTTTCTCGC AGCCATTGCC TGCCTGACCC TGAACCAGCA CCTTCTTTTC  
1314  
1315 CGAGTCATAC CCCATGATCA AAGTTTCATC GAAAACTACG CAGGGATCTT CCACTTCCAG  
1316  
1317 TTCTGGCGCT ATGGAGAGTG GGTGGACGTG GTTATAGATG ACTGCCTGCC AACGTACAAC  
1318  
1319 AATCAACTGG TTTTCACCAA GTCCAACCAC CGCAATGAGT TCTGGAGTGC TCTGCTGGAG  
1320  
1321 AAGGCTTATG CTAAGCTCCA TGGTTCCTAC GAAGCTCTGA AAGGTGGGAA CACCACAGAG  
1322  
1323 GCCATGGAGG ACTTCACAGG AGGGGTGGCA GAGTTTTTTG AGATCAGGGA TGCTCCTAGT  
1324  
1325 GACATGTACA AGATCATGAA GAAAGCCATC GAGAGAGGCT CCCTCATGGG CTGCTCCATT  
1326  
1327 GATGATGGCA CGAACATGAC CTATGGAACC TCTCCTTCTG GTCTGAACAT GGGGGAGTTG  
1328  
1329 ATTGCACGGA TGGTAAGGAA TATGGATAAC TCACTGCTCC AGGACTCAGA CCTCGACCCC  
1330  
1331 AGAGGCTCAG ATGAAAGACC GACCCGGACA ATCATTCCGG TTCAGTATGA GACAAGAATG  
1332  
1333 GCCTGCGGGC TGGTCAGAGG TCACGCCTAC TCTGTCACGG GGCTGGATGA GGTCCCCTTC  
1334  
1335 AAAGGTGAGA AAGTGAAGCT GGTGCGGCTG CGGAATCCGT GGGGCCAGGT GGAGTGGAAC  
1336  
1337 GGTTCTTGGA GTGATAGATG GAAGGACTGG AGCTTTGTGG ACAAAGATGA GAAGGCCCGT  
1338  
1339 CTGCAGCACC AGGTCACCTGA GGATGGAGAG TTCTGGATGT CCTATGAGGA TTTCATCTAC  
1340  
1341 CATTTACAAA AGTTGGAGAT CTGCAACCTC ACGGCCGATG CTCTGCAGTC TGACAAGCTT  
1342  
1343 CAGACCTGGA CAGTGTCTGT GAACGAGGGC CGCTGGGTAC GGGGTTGCTC TGCCGGAGGC  
1344  
1345 TGCCGCAACT TCCCAGATAC TTTCTGGACC AACCCTCAGT ACCGTCTGAA GCTCCTGGAG  
1346  
1347 GAGGACGATG ACCCTGATGA CTCGGAGGTG ATTTGCAGCT TCCTGGTGGC CCTGATGCAG  
1348  
1349 AAGAACCAGC GGAAGGACCG GAAGCTAGGG GCCAGTCTCT TCACCATTGG CTTCGCCATC  
1350  
1351 TACGAGGTTT CCAAAGAGAT GCACGGGAAC AAGCAGCACC TGCAGAAGGA CTTCTTCCTG  
1352  
1353 TACAACGCCT CCAAGGCCAG GAGCAAAACC TACATCAACA TGCGGGAGGT GTCCCAGCGC  
1354  
1355 TTCCGCCTGC CTCCCAGCGA GTACGTCATC GTGCCCTCCA CCTACGAGCC CCACCAGGAG  
1356  
1357 GGGGAATTCA TCCTCCGGGT CTTCTCTGAA AAGAGGAACC TCTCTGAGGA AGTTGAAAAT  
1358  
1359 ACCATCTCCG TGGATCGGCC AGTGAAAAAG AAAAAAACCA AGCCCATCAT CTTCTTTTCG  
1360  
1361 GACAGAGCAA ACAGCAACAA GGAGCTGGGT GTGGACCAGG AGTCAGAGGA GGGCAAAGGC  
1362  
1363 AAAACAAGCC CTGATAAGCA AAAGCAGTCC CCACAGCCAC AGCCTGGCAG CTCTGATCAG  
1364  
1365 GAAAGTGAGG AACAGCAACA ATTCCGGAAC ATTTTCAAGC AGATAGCAGG AGATGACATG

*same*

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:03

INPUT SET: S26725.raw

1366  
1367 GAGATCTGTG CAGATGAGCT CAAGAAGGTC CTTAACACAG TCGTGAACAA ACACAAGGAC  
1368  
1369 CTGAAGACAC ACGGGTTCAC ACTGGAGTCC TGCCGTAGCA TGATTGCGCT CATGGATACA  
1370  
1371 GATGGCTCTG GAAAGCTCAA CCTGCAGGAG TTCCACCACC TCTGGAACAA GATTAAGGCC  
1372  
1373 TGGCAGAAAA TTTTCAAACA CTATGACACA GACCAGTCCG GCACCATCAA CAGCTACGAG  
1374  
1375 ATGCGAAATG CAGTCAACGA CGCAGGATTC CACCTCAACA ACCAGCTCTA TGACATCATT  
1376  
1377 ACCATGCGGT ACGCAGACAA ACACATGAAC ATCGACTTTG ACAGTTTCAT CTGCTGCTTC  
1378  
1379 GTTAGGCTGG AGGGCATGTT CAGAGCTTTT CATGCATTTG ACAAGGATGG AGATGGTATC  
1380  
1381 ATCAAGCTCA ACGTTCTGGA GTGGCTGCAG CTCACCATGT ATGCCTGAAC CAGGCTGGCC  
1382  
1383 TCATCCAAAG CCATGCAGGA TCACTCAGGA TTTCACTTTT ACCCTCTATT TCCAAAGCCA  
1384  
1385 TTTACCTCAA AGGACCCAGC AGCTACACCC CTACAGGCTT CCAGGCACCT CATCAGTCAT  
1386  
1387 GTTCCCTCCTC CATTTTACCC CCTACCCATC CTTGATCGGT CATGCCTAGC CTGACCCCTT  
1388  
1389 AGTAAAGCAA TGAGGTAGGA AGAACAACCC CTTGTCCCTT TGCCATGTGG AGGAAAGTGC  
1390  
1391 CTGCCTCTGG TCCGAGCCGC CTCGGTCTG AAGCGAGTGC TCCTGCTTAC CTTGCTCTAG  
1392  
1393 GCTGTCTGCA GAAGCACCTG CCGGTGGCAC TCAGCACCTC CTTGTGCTAG AGCCCTCCAT  
1394  
1395 CACCTTCACG CTGTCCACC ATGGGCCAGG AACCAAACCA GCACTGGGTT CTAAGTCTGT  
1396  
1397 GGGGTAAACT AACTCAGTGG AATAGGGCTG GTTACTTTGG GCTGTCCAAC TCATAAGTTT  
1398  
1399 GGCTGCATTT TGAAAAAGC TGATCTAAAT AAAGGCATGT GTATGGCTGG TCCCCTTGTG  
1400  
1401 TTTTGTGTC TCACATTTAG ATATCAGCCA TGCATGACTG AATGGCTTCC AATCATATAC  
1402  
1403 TCACCTATCA CCTACAAGAG AACAAAGAAA AACACACACA AAAACAAAAT CTTGAATTTT  
1404  
1405 GTAATCATGC CTATTGCTAT TTCTTGAGCA TAAGAATGGC TCAGATACTT TCCAAGACAT  
1406  
1407 AAAAGGAAGG CAGAGGAATA GTTGTGCTG TAAAAGACAT CAAGAATAAA TGGGTCATGT  
1408  
1409 ACAACGGGAG GGGCCGGTTA CCTGAATAAT GGAGTGGAGA TTGAGCTATC CTAGCTCCTC  
1410  
1411 TGCTCACTAA CTGACCTGTC GCATGACCGT GGACAAAACC CTGAACGCAG CTGTTTGTGTT  
1412  
1413 GCTAACTTC TCTGGACCAT GGCCTGCGGC ATATCTATAG GCATCCTGTG TTTTCCACCC  
1414  
1415 AGTTTCCTTC TTCCTCGCTA AGCCAACGTG GAAAGGGCTG GCCGTGAATA TGCAGACAAG  
1416  
1417 GTAACGAAAG TAAACCGTCA ATTAGTAAAA GTACTTCATT TTCCTCTTGT ATTTGCTTCA  
1418

*same*

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:05

INPUT SET: S26725.raw

1419 TATCTTGCTT CACAAAGTTA CGAAGTTCAC AGCTTTATAC CAAAATGTAA GAAGGCTATT  
1420  
1421 TGCTTATAAA CATTTTGTCA GTCAGGTGTC ATCTGATTTC ATTCTTCTAA TCCATATTCA  
1422  
1423 ATATTANAAA AATCAGAAAC CAAGGGTGCT GGAGCAGCTC TAGGGCATAT ATTTCTCTTA  
1424  
1425 AATAGGAGAA AGATTTTCAA CAGCTTTTCC TCCTTGACCC CCTCCTTTCC CAATTTATTT  
1426  
1427 GGGTCACTAC CTTGAATTTA GAGTGAATCT GGGAAATGTA GTCACCAGG  
1428  
1429

*same*

---

2232 (2) INFORMATION FOR SEQ ID NO: 10:  
2233

--> 2234 (i) SEQUENCE CHARACTERISTICS:  
2235 (A) LENGTH: 20 base pairs  
2236 (B) TYPE: nucleic acid  
2237 (C) STRANDEDNESS: double  
2238 (D) TOPOLOGY: linear  
2239  
2240 (ii) MOLECULE TYPE: cDNA  
2241  
2242 (ix) FEATURE:  
2243 (A) NAME/KEY: misc\_feature  
2244 (D) OTHER INFORMATION:/label= Table 1  
2245  
2246 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:  
2247  
2248 ATGGAGCCAA CAGAACTGAC  
2249

*total*

✓

---

2250 (2) INFORMATION FOR SEQ ID NO: 11:  
2251

--> 2252 (i) SEQUENCE CHARACTERISTICS:  
2253 (A) LENGTH: 21 base pairs  
2254 (B) TYPE: nucleic acid  
2255 (C) STRANDEDNESS: double  
2256 (D) TOPOLOGY: linear  
2257  
2258 (ii) MOLECULE TYPE: Primer  
2259  
2260 (ix) FEATURE:  
2261 (A) NAME/KEY: misc\_feature  
2262 (D) OTHER INFORMATION:/label= Table 1  
2263  
2264 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:  
2265  
2266 GTATGACTCG GAAAAGAAGGT  
2267

✓

---

2268 (2) INFORMATION FOR SEQ ID NO: 12:  
2269

2270 (i) SEQUENCE CHARACTERISTICS:

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:06

INPUT SET: S26725.raw

--> 2271 (A) LENGTH: 20 base pairs  
2272 (B) TYPE: nucleic acid  
2273 (C) STRANDEDNESS: double  
2274 (D) TOPOLOGY: linear  
2275  
2276 (ii) MOLECULE TYPE: Primer  
2277  
2278 (ix) FEATURE:  
2279 (A) NAME/KEY: misc\_feature  
2280 (D) OTHER INFORMATION:/label= Table 1  
2281  
2282 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:  
2283  
2284 TAAGCAAAAG CAGTCCCCAC  
2285

total  
✓

---

2286 (2) INFORMATION FOR SEQ ID NO: 13:  
2287  
2288 (i) SEQUENCE CHARACTERISTICS:  
--> 2289 (A) LENGTH: 21 base pairs  
2290 (B) TYPE: nucleic acid  
2291 (C) STRANDEDNESS: double  
2292 (D) TOPOLOGY: linear  
2293  
2294 (ii) MOLECULE TYPE: primer  
2295  
2296 (ix) FEATURE:  
2297 (A) NAME/KEY: misc\_feature  
2298 (D) OTHER INFORMATION:/label= Table 1  
2299  
2300 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:  
2301  
2302 TTGCTGTTCC TCACTTTCCTG  
2303

✓

---

2304 (2) INFORMATION FOR SEQ ID NO: 14:  
2305  
2306 (i) SEQUENCE CHARACTERISTICS:  
--> 2307 (A) LENGTH: 20 base pairs  
2308 (B) TYPE: nucleic acid  
2309 (C) STRANDEDNESS: double  
2310 (D) TOPOLOGY: linear  
2311  
2312 (ii) MOLECULE TYPE: primer  
2313  
2314 (ix) FEATURE:  
2315 (A) NAME/KEY: misc\_feature  
2316 (D) OTHER INFORMATION:/label= Table 1  
2317  
2318 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:  
2319  
2320 GTTTCATCTG CTGCTTCGTT  
2321

✓

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:08

INPUT SET: S26725.raw

---

2322 (2) INFORMATION FOR SEQ ID NO: 15:  
2323  
2324 (i) SEQUENCE CHARACTERISTICS:  
--> 2325 (A) LENGTH: 20 base pairs  
2326 (B) TYPE: nucleic acid  
2327 (C) STRANDEDNESS: double  
2328 (D) TOPOLOGY: linear  
2329  
2330 (ii) MOLECULE TYPE: primer  
2331  
2332 (ix) FEATURE:  
2333 (A) NAME/KEY: misc\_feature  
2334 (D) OTHER INFORMATION:/label= Table 1  
2335  
2336 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:  
2337  
2338 CTGGTTCAGG CATA CATGGT  
2339

total

✓

---

2340 (2) INFORMATION FOR SEQ ID NO: 16:  
2341  
2342 (i) SEQUENCE CHARACTERISTICS:  
--> 2343 (A) LENGTH: 22 base pairs  
2344 (B) TYPE: nucleic acid  
2345 (C) STRANDEDNESS: double  
2346 (D) TOPOLOGY: linear  
2347  
2348 (ii) MOLECULE TYPE: primer  
2349  
2350 (ix) FEATURE:  
2351 (A) NAME/KEY: misc\_feature  
2352 (D) OTHER INFORMATION:/label= Table 1  
2353  
2354 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:  
2355  
2356  
2357 TTCTTTATGT GGACCCTGAG TT  
2358

✓

---

2359 (2) INFORMATION FOR SEQ ID NO: 17:  
2360  
2361 (i) SEQUENCE CHARACTERISTICS:  
--> 2362 (A) LENGTH: 19 base pairs  
2363 (B) TYPE: nucleic acid  
2364 (C) STRANDEDNESS: double  
2365 (D) TOPOLOGY: linear  
2366  
2367 (ii) MOLECULE TYPE: primer  
2368  
2369 (ix) FEATURE:  
2370 (A) NAME/KEY: misc\_feature  
2371 (D) OTHER INFORMATION:/label= Table 1

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:09

INPUT SET: S26725.raw

2372  
2373 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:  
2374  
2375 ACGAACTGGA TGGGGAAC  
2376

total  
✓

2377 (2) INFORMATION FOR SEQ ID NO:18 :  
2378  
2379 (i) SEQUENCE CHARACTERISTICS:  
--> 2380 (A) LENGTH: 20 base pairs  
2381 (B) TYPE: nucleic acid  
2382 (C) STRANDEDNESS: double  
2383 (D) TOPOLOGY: linear  
2384  
2385 (ii) MOLECULE TYPE: primer  
2386  
2387 (ix) FEATURE:  
2388 (A) NAME/KEY: misc\_feature  
2389 (D) OTHER INFORMATION:/label= Table 3  
2390  
2391 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:  
2392  
2393 TTCAGTACCTCCCGTTCACC  
2394

invalid - put 1.822 of sequence into  
group bases (non-coding) into 10's

total  
✓

2395 (2) INFORMATION FOR SEQ ID NO:19:  
2396  
2397 (i) SEQUENCE CHARACTERISTICS:  
--> 2398 (A) LENGTH: 20 base pairs  
2399 (B) TYPE: nucleic acid  
2400 (C) STRANDEDNESS: double  
2401 (D) TOPOLOGY: linear  
2402  
2403 (ii) MOLECULE TYPE: primer  
2404  
2405 (ix) FEATURE:  
2406 (A) NAME/KEY: misc\_feature  
2407 (D) OTHER INFORMATION:/label= Table 3  
2408  
2409 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:  
2410  
2411 GATGCTTGAGCCAGGAAAAC  
2412

This type of error  
appears in all  
subsequent nucleic  
acid sequences

✓

2413 (2) INFORMATION FOR SEQ ID NO:20:  
2414  
2415 (i) SEQUENCE CHARACTERISTICS:  
--> 2416 (A) LENGTH: 22 base pairs  
2417 (B) TYPE: nucleic acid  
2418 (C) STRANDEDNESS: double  
2419 (D) TOPOLOGY: linear  
2420  
2421 (ii) MOLECULE TYPE: primer



RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:11

INPUT SET: S26725.raw

2422  
2423 (ix) FEATURE:  
2424 (A) NAME/KEY: misc\_feature  
2425 (D) OTHER INFORMATION:/label= Table 3  
2426  
2427 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:  
2428  
2429 CTTTCCTTGAAGGTAGCTGTAT  
2430

2431 (2) INFORMATION FOR SEQ ID NO:21 :  
2432  
2433 (i) SEQUENCE CHARACTERISTICS:  
--> 2434 (A) LENGTH: 20 base pairs  
2435 (B) TYPE: nucleic acid  
2436 (C) STRANDEDNESS: double  
2437 (D) TOPOLOGY: linear  
2438  
2439 (ii) MOLECULE TYPE: primer  
2440  
2441 (ix) FEATURE:  
2442 (A) NAME/KEY: misc\_feature  
2443 (D) OTHER INFORMATION:/label= Table 3  
2444  
2445 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:21 :  
2446  
2447 GAGGTGCTGAGTGAGAGGAC  
2448

2449 (2) INFORMATION FOR SEQ ID NO:22 :  
2450  
2451 (i) SEQUENCE CHARACTERISTICS:  
--> 2452 (A) LENGTH: 22 base pairs  
2453 (B) TYPE: nucleic acid  
2454 (C) STRANDEDNESS: double  
2455 (D) TOPOLOGY: linear  
2456  
2457 (ii) MOLECULE TYPE: primer  
2458  
2459 (ix) FEATURE:  
2460 (A) NAME/KEY: misc\_feature  
2461 (D) OTHER INFORMATION:/label= Table 3  
2462  
2463 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22 :  
2464  
2465 ACTCCGTCTCAAAAAATACCT  
2466

2467 (2) INFORMATION FOR SEQ ID NO:23:  
2468  
2469 (i) SEQUENCE CHARACTERISTICS:  
--> 2470 (A) LENGTH: 20 base pairs  
2471 (B) TYPE: nucleic acid

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:12

INPUT SET: S26725.raw

2472 (C) STRANDEDNESS: double  
2473 (D) TOPOLOGY: linear  
2474  
2475 (ii) MOLECULE TYPE: primer  
2476  
2477 (ix) FEATURE:  
2478 (A) NAME/KEY: misc\_feature  
2479 (D) OTHER INFORMATION:/label= Table 3  
2480  
2481 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23 :  
2482  
2483 ATTGTCCTTTACCTCTGG  
2484

2485 (2) INFORMATION FOR SEQ ID NO:24 :  
2486  
2487 (i) SEQUENCE CHARACTERISTICS:  
--> 2488 (A) LENGTH: 20 base pairs  
2489 (B) TYPE: nucleic acid  
2490 (C) STRANDEDNESS: double  
2491 (D) TOPOLOGY: linear  
2492  
2493 (ii) MOLECULE TYPE: primer  
2494  
2495 (ix) FEATURE:  
2496 (A) NAME/KEY: misc\_feature  
2497 (D) OTHER INFORMATION:/label= Table 3  
2498  
2499 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:24 :  
2500  
2501 TGGAAGTAGGAGAGTGGGCA  
2502

2503 (2) INFORMATION FOR SEQ ID NO:25 :  
2504  
2505 (i) SEQUENCE CHARACTERISTICS:  
--> 2506 (A) LENGTH: 20 base pairs  
2507 (B) TYPE: nucleic acid  
2508 (C) STRANDEDNESS: double  
2509 (D) TOPOLOGY: linear  
2510  
2511 (ii) MOLECULE TYPE: primer  
2512  
2513 (ix) FEATURE:  
2514 (A) NAME/KEY: misc\_feature  
2515 (D) OTHER INFORMATION:/label= Table 3  
2516  
2517 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:25 :  
2518  
2519 GGGTAGATGGGTGGGAAGTT  
2520

2521 (2) INFORMATION FOR SEQ ID NO:26 :

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:14

INPUT SET: S26725.raw

2522  
2523 (i) SEQUENCE CHARACTERISTICS:  
--> 2524 (A) LENGTH: 20 base pairs  
2525 (B) TYPE: nucleic acid  
2526 (C) STRANDEDNESS: double  
2527 (D) TOPOLOGY: linear  
2528  
2529 (ii) MOLECULE TYPE: primer  
2530  
2531 (ix) FEATURE:  
2532 (A) NAME/KEY: misc\_feature  
2533 (D) OTHER INFORMATION:/label= Table 3  
2534  
2535 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:26 :  
2536  
2537 GAGGAATGTGGAGGAAGGAC  
2538

---

2539 (2) INFORMATION FOR SEQ ID NO:27:  
2540  
2541 (i) SEQUENCE CHARACTERISTICS:  
--> 2542 (A) LENGTH: 20 base pairs  
2543 (B) TYPE: nucleic acid  
2544 (C) STRANDEDNESS: double  
2545 (D) TOPOLOGY: linear  
2546  
2547 (ii) MOLECULE TYPE: primer  
2548  
2549 (ix) FEATURE:  
2550 (A) NAME/KEY: misc\_feature  
2551 (D) OTHER INFORMATION:/label= Table 3  
2552  
2553 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:  
2554  
2555 TTCCTGTGAGTGAGGTCTCG  
2556

---

2557 (2) INFORMATION FOR SEQ ID NO:28 :  
2558  
2559 (i) SEQUENCE CHARACTERISTICS:  
--> 2560 (A) LENGTH: 20 base pairs  
2561 (B) TYPE: nucleic acid  
2562 (C) STRANDEDNESS: double  
2563 (D) TOPOLOGY: linear  
2564  
2565 (ii) MOLECULE TYPE: primer  
2566  
2567 (ix) FEATURE:  
2568 (A) NAME/KEY: misc\_feature  
2569 (D) OTHER INFORMATION:/label= Table 3  
2570  
2571 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:28 :  
2572

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:16

INPUT SET: S26725.raw

2573 GGAACTCTGTGACCCCAAAT  
25742575 (2) INFORMATION FOR SEQ ID NO:29 :  
2576

--> 2577 (i) SEQUENCE CHARACTERISTICS:  
2578 (A) LENGTH: 20 base pairs  
2579 (B) TYPE: nucleic acid  
2580 (C) STRANDEDNESS: double  
2581 (D) TOPOLOGY: linear  
2582  
2583 (ii) MOLECULE TYPE: primer  
2584  
2585 (ix) FEATURE:  
2586 (A) NAME/KEY: misc\_feature  
2587 (D) OTHER INFORMATION:/label= Table 3  
2588  
2589 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:29 :  
2590  
2591 TCCTCAAACAAAACATTTCGC  
2592

2593 (2) INFORMATION FOR SEQ ID NO:30 :  
2594

--> 2595 (i) SEQUENCE CHARACTERISTICS:  
2596 (A) LENGTH: 20 base pairs  
2597 (B) TYPE: nucleic acid  
2598 (C) STRANDEDNESS: double  
2599 (D) TOPOLOGY: linear  
2600  
2601 (ii) MOLECULE TYPE: primer  
2602  
2603 (ix) FEATURE:  
2604 (A) NAME/KEY: misc\_feature  
2605 (D) OTHER INFORMATION:/label= Table 3  
2606  
2607 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:  
2608  
2609 GTTCCCTACATTCTCCATCG  
2610

2611 (2) INFORMATION FOR SEQ ID NO:31 :  
2612

--> 2613 (i) SEQUENCE CHARACTERISTICS:  
2614 (A) LENGTH: 21 base pairs  
2615 (B) TYPE: nucleic acid  
2616 (C) STRANDEDNESS: double  
2617 (D) TOPOLOGY: linear  
2618  
2619 (ii) MOLECULE TYPE: primer  
2620  
2621 (ix) FEATURE:  
2622 (A) NAME/KEY: misc\_feature

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:17

INPUT SET: S26725.raw

2623 (D) OTHER INFORMATION:/label= Table 3

2624

2625 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:31 :

2626

2627 GTTATTTCAACCCAGACCCTT

2628

2629 (2) INFORMATION FOR SEQ ID NO:32 :

2630

(i) SEQUENCE CHARACTERISTICS:

--&gt; 2632 (A) LENGTH: 21 base pairs

2633 (B) TYPE: nucleic acid

2634 (C) STRANDEDNESS: double

2635 (D) TOPOLOGY: linear

2636

2637 (ii) MOLECULE TYPE: primer

2638

2639 (ix) FEATURE:

2640 (A) NAME/KEY: misc\_feature

2641 (D) OTHER INFORMATION:/label= Table 3

2642

2643 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:32 :

2644

2645 AATGGGTTCTCTGGTTACTGC

2646

2647 (2) INFORMATION FOR SEQ ID NO:33 :

2648

(i) SEQUENCE CHARACTERISTICS:

--&gt; 2650 (A) LENGTH: 21 base pairs

2651 (B) TYPE: nucleic acid

2652 (C) STRANDEDNESS: double

2653 (D) TOPOLOGY: linear

2654

2655 (ii) MOLECULE TYPE: primer

2656

(ix) FEATURE:

2657 (A) NAME/KEY: misc\_feature

2658 (D) OTHER INFORMATION:/label= Table 3

2659

2660 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:33 :

2661

2662 AGCACGAAAAGCAAAGATAAA

2663

2665 (2) INFORMATION FOR SEQ ID NO:34 :

2666

(i) SEQUENCE CHARACTERISTICS:

--&gt; 2668 (A) LENGTH: 20 base pairs

2669 (B) TYPE: nucleic acid

2670 (C) STRANDEDNESS: double

2671 (D) TOPOLOGY: linear

2672

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:19

INPUT SET: S26725.raw

2673 (ii) MOLECULE TYPE: primer  
2674  
2675 (ix) FEATURE:  
2676 (A) NAME/KEY: misc\_feature  
2677 (D) OTHER INFORMATION:/label= Table 3  
2678  
2679 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:34 :  
2680  
2681 GTAAGAGATTGCCCCCAG  
2682

---

2683 (2) INFORMATION FOR SEQ ID NO:35 :  
2684  
2685 (i) SEQUENCE CHARACTERISTICS:  
--> 2686 (A) LENGTH: 20 base pairs  
2687 (B) TYPE: nucleic acid  
2688 (C) STRANDEDNESS: double  
2689 (D) TOPOLOGY: linear  
2690  
2691 (ii) MOLECULE TYPE: primer  
2692  
2693 (ix) FEATURE:  
2694 (A) NAME/KEY: misc\_feature  
2695 (D) OTHER INFORMATION:/label= Table 3  
2696  
2697 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:35 :  
2698  
2699 TCTGCGGATCATTGTTTGT  
2700

---

2701 (2) INFORMATION FOR SEQ ID NO:36 :  
2702  
2703 (i) SEQUENCE CHARACTERISTICS:  
--> 2704 (A) LENGTH: 20 base pairs  
2705 (B) TYPE: nucleic acid  
2706 (C) STRANDEDNESS: double  
2707 (D) TOPOLOGY: linear  
2708  
2709 (ii) MOLECULE TYPE: primer  
2710  
2711 (ix) FEATURE:  
2712 (A) NAME/KEY: misc\_feature  
2713 (D) OTHER INFORMATION:/label= Table 3  
2714  
2715 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:36 :  
2716  
2717 CCTTCCCTTCTTCCTGCTTC  
2718

---

2719 (2) INFORMATION FOR SEQ ID NO:37 :  
2720  
2721 (i) SEQUENCE CHARACTERISTICS:  
--> 2722 (A) LENGTH: 20 base pairs

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:20

INPUT SET: S26725.raw

2723 (B) TYPE: nucleic acid  
2724 (C) STRANDEDNESS: double  
2725 (D) TOPOLOGY: linear  
2726  
2727 (ii) MOLECULE TYPE: primer  
2728  
2729 (ix) FEATURE:  
2730 (A) NAME/KEY: misc\_feature  
2731 (D) OTHER INFORMATION:/label= Table 3  
2732  
2733 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:37 :  
2734  
2735 CTCTCTTCCCCACCCCTTACC  
2736

---

2737 (2) INFORMATION FOR SEQ ID NO:38 :  
2738  
2739 (i) SEQUENCE CHARACTERISTICS:  
--> 2740 (A) LENGTH: 20 base pairs  
2741 (B) TYPE: nucleic acid  
2742 (C) STRANDEDNESS: double  
2743 (D) TOPOLOGY: linear  
2744  
2745 (ii) MOLECULE TYPE: primer  
2746  
2747 (ix) FEATURE:  
2748 (A) NAME/KEY: misc\_feature  
2749 (D) OTHER INFORMATION:/label= Table 3  
2750  
2751 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:38 :  
2752  
2753 CCTCCTCACCTGCTCCCATA  
2754

---

2755 (2) INFORMATION FOR SEQ ID NO:39 :  
2756  
2757 (i) SEQUENCE CHARACTERISTICS:  
--> 2758 (A) LENGTH: 20 base pairs  
2759 (B) TYPE: nucleic acid  
2760 (C) STRANDEDNESS: double  
2761 (D) TOPOLOGY: linear  
2762  
2763 (ii) MOLECULE TYPE: primer  
2764  
2765 (ix) FEATURE:  
2766 (A) NAME/KEY: misc\_feature  
2767 (D) OTHER INFORMATION:/label= Table 3  
2768  
2769 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:39 :  
2770  
2771 TTTTTCGGCTTAGACCCCTCC  
2772

---

2773 (2) INFORMATION FOR SEQ ID NO:40 :

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:22

INPUT SET: S26725.raw

2774  
2775 (i) SEQUENCE CHARACTERISTICS:  
--> 2776 (A) LENGTH: 22 base pairs  
2777 (B) TYPE: nucleic acid  
2778 (C) STRANDEDNESS: double  
2779 (D) TOPOLOGY: linear  
2780  
2781 (ii) MOLECULE TYPE: primer  
2782  
2783 (ix) FEATURE:  
2784 (A) NAME/KEY: misc\_feature  
2785 (D) OTHER INFORMATION:/label= Table 3  
2786  
2787 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:40 :  
2788  
2789 TGTGGGGAATAGAAATAAATGG  
2790

---

2791 (2) INFORMATION FOR SEQ ID NO:41 :  
2792  
2793 (i) SEQUENCE CHARACTERISTICS:  
--> 2794 (A) LENGTH: 19 base pairs  
2795 (B) TYPE: nucleic acid  
2796 (C) STRANDEDNESS: double  
2797 (D) TOPOLOGY: linear  
2798  
2799 (ii) MOLECULE TYPE: primer  
2800  
2801 (ix) FEATURE:  
2802 (A) NAME/KEY: misc\_feature  
2803 (D) OTHER INFORMATION:/label= Table 3  
2804  
2805 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:41 :  
2806  
2807 CCAGGAGCTCTGTGGGTCA  
2808

---

2809 (2) INFORMATION FOR SEQ ID NO:42 :  
2810  
2811 (i) SEQUENCE CHARACTERISTICS:  
--> 2812 (A) LENGTH: 21 base pairs  
2813 (B) TYPE: nucleic acid  
2814 (C) STRANDEDNESS: double  
2815 (D) TOPOLOGY: linear  
2816  
2817 (ii) MOLECULE TYPE: primer  
2818  
2819 (ix) FEATURE:  
2820 (A) NAME/KEY: misc\_feature  
2821 (D) OTHER INFORMATION:/label= Table 3  
2822  
2823 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:42 :  
2824



RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:23

INPUT SET: S26725.fw

2825 GGCTCCTCATCCTCATTACA  
28262827 (2) INFORMATION FOR SEQ ID NO:43 :  
28282829 (i) SEQUENCE CHARACTERISTICS:  
--> 2830 (A) LENGTH: 20 base pairs  
2831 (B) TYPE: nucleic acid  
2832 (C) STRANDEDNESS: double  
2833 (D) TOPOLOGY: linear  
28342835 (ii) MOLECULE TYPE: primer  
2836

2837 (ix) FEATURE:

2838 (A) NAME/KEY: misc\_feature  
2839 (D) OTHER INFORMATION:/label= Table 3  
28402841 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:43:  
28422843 GTGGAGGAGGGTGAGTGTGC  
28442845 (2) INFORMATION FOR SEQ ID NO:44 :  
28462847 (i) SEQUENCE CHARACTERISTICS:  
--> 2848 (A) LENGTH: 20 base pairs  
2849 (B) TYPE: nucleic acid  
2850 (C) STRANDEDNESS: double  
2851 (D) TOPOLOGY: linear  
28522853 (ii) MOLECULE TYPE: primer  
2854

2855 (ix) FEATURE:

2856 (A) NAME/KEY: misc\_feature  
2857 (D) OTHER INFORMATION:/label= Table 3  
28582859 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:44 :  
28602861 TGTGGCAGGACAGGACGTTC  
28622863 (2) INFORMATION FOR SEQ ID NO:45 :  
28642865 (i) SEQUENCE CHARACTERISTICS:  
--> 2866 (A) LENGTH: 20 base pairs  
2867 (B) TYPE: nucleic acid  
2868 (C) STRANDEDNESS: double  
2869 (D) TOPOLOGY: linear  
28702871 (ii) MOLECULE TYPE: primer  
2872

2873 (ix) FEATURE:

2874 (A) NAME/KEY: misc\_feature

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:25

INPUT SET: S26725.raw

2875 (D) OTHER INFORMATION:/label= Table 3  
2876  
2877 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:45 :  
2878  
2879 TTCAACCTCTGGAGTGGGCC  
2880

↓

2881 (2) INFORMATION FOR SEQ ID NO:46 :  
2882  
2883 (i) SEQUENCE CHARACTERISTICS:  
--> 2884 (A) LENGTH: 20 base pairs  
2885 (B) TYPE: nucleic acid  
2886 (C) STRANDEDNESS: double  
2887 (D) TOPOLOGY: linear  
2888  
2889 (ii) MOLECULE TYPE: primer  
2890  
2891 (ix) FEATURE:  
2892 (A) NAME/KEY: misc\_feature  
2893 (D) OTHER INFORMATION:/label= Table 3  
2894  
2895 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:46 :  
2896  
2897 CACCAGAGCAAACCGTCCAC  
2898

↓

2899 (2) INFORMATION FOR SEQ ID NO:47 :  
2900  
2901 (i) SEQUENCE CHARACTERISTICS:  
--> 2902 (A) LENGTH: 20 base pairs  
2903 (B) TYPE: nucleic acid  
2904 (C) STRANDEDNESS: double  
2905 (D) TOPOLOGY: linear  
2906  
2907 (ii) MOLECULE TYPE: primer  
2908  
2909 (ix) FEATURE:  
2910 (A) NAME/KEY: misc\_feature  
2911 (D) OTHER INFORMATION:/label= Table 3  
2912  
2913 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:47 :  
2914  
2915 ACAGCCAGACTCCCATTCC  
2916

↓

2917 (2) INFORMATION FOR SEQ ID NO:48 :  
2918  
2919 (i) SEQUENCE CHARACTERISTICS:  
--> 2920 (A) LENGTH: 20 base pairs  
2921 (B) TYPE: nucleic acid  
2922 (C) STRANDEDNESS: double  
2923 (D) TOPOLOGY: linear  
2924

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:26

INPUT SET: S26725.raw

2925 (ii) MOLECULE TYPE: primer  
2926  
2927 (ix) FEATURE:  
2928 (A) NAME/KEY: misc\_feature  
2929 (D) OTHER INFORMATION:/label= Table 3  
2930  
2931 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:48 :  
2932  
2933 TTCTCTTCTCCCTTCACCTT  
2934

---

2935 (2) INFORMATION FOR SEQ ID NO:49 :  
2936  
2937 (i) SEQUENCE CHARACTERISTICS:  
--> 2938 (A) LENGTH: 22 base pairs  
2939 (B) TYPE: nucleic acid  
2940 (C) STRANDEDNESS: double  
2941 (D) TOPOLOGY: linear  
2942  
2943 (ii) MOLECULE TYPE: primer  
2944  
2945 (ix) FEATURE:  
2946 (A) NAME/KEY: misc\_feature  
2947 (D) OTHER INFORMATION:/label= Table 3  
2948  
2949 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:49 :  
2950  
2951 ACACACTTCATGCTCTCTACCC  
2952

---

2953 (2) INFORMATION FOR SEQ ID NO:50:  
2954  
2955 (i) SEQUENCE CHARACTERISTICS:  
--> 2956 (A) LENGTH: 20 base pairs  
2957 (B) TYPE: nucleic acid  
2958 (C) STRANDEDNESS: double  
2959 (D) TOPOLOGY: linear  
2960  
2961 (ii) MOLECULE TYPE: primer  
2962  
2963 (ix) FEATURE:  
2964 (A) NAME/KEY: misc\_feature  
2965 (D) OTHER INFORMATION:/label= Table 3  
2966  
2967 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:50 :  
2968  
2969 CCGCCTATTCCTTTCTCTT  
2970

---

2971 (2) INFORMATION FOR SEQ ID NO:51 :  
2972  
2973 (i) SEQUENCE CHARACTERISTICS:  
--> 2974 (A) LENGTH: 20 base pairs

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:28

INPUT SET: S26725.raw

2975 (B) TYPE: nucleic acid  
2976 (C) STRANDEDNESS: double  
2977 (D) TOPOLOGY: linear  
2978  
2979 (ii) MOLECULE TYPE: primer  
2980  
2981 (ix) FEATURE:  
2982 (A) NAME/KEY: misc\_feature  
2983 (D) OTHER INFORMATION:/label= Table 3  
2984  
2985 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:51 :  
2986  
2987 GACAAACTCCTGGGAAGCCT  
2988

✓

2989 (2) INFORMATION FOR SEQ ID NO:52 :  
2990  
2991 (i) SEQUENCE CHARACTERISTICS:  
--> 2992 (A) LENGTH: 20 base pairs  
2993 (B) TYPE: nucleic acid  
2994 (C) STRANDEDNESS: double  
2995 (D) TOPOLOGY: linear  
2996  
2997 (ii) MOLECULE TYPE: primer  
2998  
2999 (ix) FEATURE:  
3000 (A) NAME/KEY: misc\_feature  
3001 (D) OTHER INFORMATION:/label= Table 3  
3002  
3003 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:52 :  
3004  
3005 ACCTCTGACCCCTGTGAACC  
3006

✓

3007 (2) INFORMATION FOR SEQ ID NO:53:  
3008  
3009 (i) SEQUENCE CHARACTERISTICS:  
--> 3010 (A) LENGTH: 20 base pairs  
3011 (B) TYPE: nucleic acid  
3012 (C) STRANDEDNESS: double  
3013 (D) TOPOLOGY: linear  
3014  
3015 (ii) MOLECULE TYPE: primer  
3016  
3017 (ix) FEATURE:  
3018 (A) NAME/KEY: misc\_feature  
3019 (D) OTHER INFORMATION:/label= Table 3  
3020  
3021 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:53 :  
3022  
3023 TGTGGATTTGTGTGCTACGC  
3024

✓

3025 (2) INFORMATION FOR SEQ ID NO:54 :

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:30

INPUT SET: S26725.raw

3026  
3027 (i) SEQUENCE CHARACTERISTICS:  
--> 3028 (A) LENGTH: 21 base pairs  
3029 (B) TYPE: nucleic acid  
3030 (C) STRANDEDNESS: double  
3031 (D) TOPOLOGY: linear  
3032  
3033 (ii) MOLECULE TYPE: primer  
3034  
3035 (ix) FEATURE:  
3036 (A) NAME/KEY: misc\_feature  
3037 (D) OTHER INFORMATION:/label= Table 3  
3038  
3039 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:54 :  
3040  
3041 CATAAATAGCACCGACAGGGA  
3042

✓

---

3043 (2) INFORMATION FOR SEQ ID NO:55 :  
3044  
3045 (i) SEQUENCE CHARACTERISTICS:  
--> 3046 (A) LENGTH: 20 base pairs  
3047 (B) TYPE: nucleic acid  
3048 (C) STRANDEDNESS: double  
3049 (D) TOPOLOGY: linear  
3050  
3051 (ii) MOLECULE TYPE: primer  
3052  
3053 (ix) FEATURE:  
3054 (A) NAME/KEY: misc\_feature  
3055 (D) OTHER INFORMATION:/label= Table 3  
3056  
3057 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:55 :  
3058  
3059 GGGATGGAGAAGAGTGAGGA  
3060

✓

---

3061 (2) INFORMATION FOR SEQ ID NO:56 :  
3062  
3063 (i) SEQUENCE CHARACTERISTICS:  
--> 3064 (A) LENGTH: 20 base pairs  
3065 (B) TYPE: nucleic acid  
3066 (C) STRANDEDNESS: double  
3067 (D) TOPOLOGY: linear  
3068  
3069 (ii) MOLECULE TYPE: primer  
3070  
3071 (ix) FEATURE:  
3072 (A) NAME/KEY: misc\_feature  
3073 (D) OTHER INFORMATION:/label= Table 3  
3074  
3075 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:56 :  
3076

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:31INPUT SET: S26725.raw  
↓3077 TCCTCACTCTTCTCCATCCC  
30783079 (2) INFORMATION FOR SEQ ID NO:57 :  
3080

--> 3081 (i) SEQUENCE CHARACTERISTICS:  
3082 (A) LENGTH: 19 base pairs  
3083 (B) TYPE: nucleic acid  
3084 (C) STRANDEDNESS: double  
3085 (D) TOPOLOGY: linear  
3086

3087 (ii) MOLECULE TYPE: primer  
3088

3089 (ix) FEATURE:  
3090 (A) NAME/KEY: misc\_feature  
3091 (D) OTHER INFORMATION:/label= Table 3  
3092

3093 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:57 :  
30943095 ACCCTGATGTTGCCTTGG  
3096 ✓3097 (2) INFORMATION FOR SEQ ID NO:58 :  
3098

--> 3099 (i) SEQUENCE CHARACTERISTICS:  
3100 (A) LENGTH: 20 base pairs  
3101 (B) TYPE: nucleic acid  
3102 (C) STRANDEDNESS: double  
3103 (D) TOPOLOGY: linear  
3104

3105 (ii) MOLECULE TYPE: primer  
3106

3107 (ix) FEATURE:  
3108 (A) NAME/KEY: misc\_feature  
3109 (D) OTHER INFORMATION:/label= Table 3  
3110

3111 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:58 :  
31123113 GGGGATTTTGCTGTGTGCTG  
3114 ✓3115 (2) INFORMATION FOR SEQ ID NO:59 :  
3116

--> 3117 (i) SEQUENCE CHARACTERISTICS:  
3118 (A) LENGTH: 20 base pairs  
3119 (B) TYPE: nucleic acid  
3120 (C) STRANDEDNESS: double  
3121 (D) TOPOLOGY: linear  
3122

3123 (ii) MOLECULE TYPE: primer  
3124

3125 (ix) FEATURE:  
3126 (A) NAME/KEY: misc\_feature

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:33

INPUT SET: S26725.raw

3127 (D) OTHER INFORMATION:/label= Table 3

3128

3129 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:59 :

3130

3131 ATTCCTGCTCCCACCGTCTC

3132

3133 (2) INFORMATION FOR SEQ ID NO:60:

3134

3135 (i) SEQUENCE CHARACTERISTICS:

--&gt; 3136 (A) LENGTH: 20 base pairs

3137 (B) TYPE: nucleic acid

3138 (C) STRANDEDNESS: double

3139 (D) TOPOLOGY: linear

3140

3141 (ii) MOLECULE TYPE: primer

3142

3143 (ix) FEATURE:

3144 (A) NAME/KEY: misc\_feature

3145 (D) OTHER INFORMATION:/label= Table 3

3146

3147 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:60 :

3148

3149 CACAGAGTGTCCGAGAGGCA

3150

3151 (2) INFORMATION FOR SEQ ID NO:61 :

3152

3153 (i) SEQUENCE CHARACTERISTICS:

--&gt; 3154 (A) LENGTH: 22 base pairs

3155 (B) TYPE: nucleic acid

3156 (C) STRANDEDNESS: double

3157 (D) TOPOLOGY: linear

3158

3159 (ii) MOLECULE TYPE: primer

3160

3161 (ix) FEATURE:

3162 (A) NAME/KEY: misc\_feature

3163 (D) OTHER INFORMATION:/label= Table 3

3164

3165 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:61 :

3166

3167 GGAGATTATCAGGTGAGATGCC

3168

3169 (2) INFORMATION FOR SEQ ID NO:62 :

3170

3171 (i) SEQUENCE CHARACTERISTICS:

--&gt; 3172 (A) LENGTH: 21 base pairs

3173 (B) TYPE: nucleic acid

3174 (C) STRANDEDNESS: double

3175 (D) TOPOLOGY: linear

3176

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:34

INPUT SET: S26725.raw

3177 (ii) MOLECULE TYPE: primer  
3178  
3179 (ix) FEATURE:  
3180 (A) NAME/KEY: misc\_feature  
3181 (D) OTHER INFORMATION:/label= Table 3  
3182  
3183 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:62 :  
3184  
3185 CAGAGTGTCCGAGAGGCAGGG  
3186

✓

3187 (2) INFORMATION FOR SEQ ID NO:63 :  
3188  
3189 (i) SEQUENCE CHARACTERISTICS:  
--> 3190 (A) LENGTH: 20 base pairs  
3191 (B) TYPE: nucleic acid  
3192 (C) STRANDEDNESS: double  
3193 (D) TOPOLOGY: linear  
3194  
3195 (ii) MOLECULE TYPE: primer  
3196  
3197 (ix) FEATURE:  
3198 (A) NAME/KEY: misc\_feature  
3199 (D) OTHER INFORMATION:/label= Table 3  
3200  
3201 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:63 :  
3202  
3203 CGTTGACCCCTCCACCTTGA  
3204

✓

3205 (2) INFORMATION FOR SEQ ID NO:64 :  
3206  
3207 (i) SEQUENCE CHARACTERISTICS:  
--> 3208 (A) LENGTH: 20 base pairs  
3209 (B) TYPE: nucleic acid  
3210 (C) STRANDEDNESS: double  
3211 (D) TOPOLOGY: linear  
3212  
3213 (ii) MOLECULE TYPE: primer  
3214  
3215 (ix) FEATURE:  
3216 (A) NAME/KEY: misc\_feature  
3217 (D) OTHER INFORMATION:/label= Table 3  
3218  
3219 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:64 :  
3220  
3221 GGGAAAACATGCACCTTCTT  
3222

✓

3223 (2) INFORMATION FOR SEQ ID NO:65:  
3224  
3225 (i) SEQUENCE CHARACTERISTICS:  
--> 3226 (A) LENGTH: 20 base pairs



RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/836,734DATE: 06/16/98  
TIME: 13:37:36

INPUT SET: S26725.raw

3227 (B) TYPE: nucleic acid  
3228 (C) STRANDEDNESS: double  
3229 (D) TOPOLOGY: linear  
3230  
3231 (ii) MOLECULE TYPE: primer  
3232  
3233 (ix) FEATURE:  
3234 (A) NAME/KEY: misc\_feature  
3235 (D) OTHER INFORMATION:/label= Table 3  
3236  
3237 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:65 :  
3238  
3239 TAGGGGGTAAAATGGAGGAG  
3240

---

3241 (2) INFORMATION FOR SEQ ID NO:66 :  
3242  
3243 (i) SEQUENCE CHARACTERISTICS:  
--> 3244 (A) LENGTH: 20 base pairs  
3245 (B) TYPE: nucleic acid  
3246 (C) STRANDEDNESS: double  
3247 (D) TOPOLOGY: linear  
3248  
3249 (ii) MOLECULE TYPE: primer  
3250  
3251 (ix) FEATURE:  
3252 (A) NAME/KEY: misc\_feature  
3253 (D) OTHER INFORMATION:/label= Table 3  
3254  
3255 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:66 :  
3256  
3257 ACTAACTCAGTGAATAGGG  
3258

---

3259 (2) INFORMATION FOR SEQ ID NO:67 :  
3260  
3261 (i) SEQUENCE CHARACTERISTICS:  
--> 3262 (A) LENGTH: 20 base pairs  
3263 (B) TYPE: nucleic acid  
3264 (C) STRANDEDNESS: double  
3265 (D) TOPOLOGY: linear  
3266  
3267 (ii) MOLECULE TYPE: primer  
3268  
3269 (ix) FEATURE:  
3270 (A) NAME/KEY: misc\_feature  
3271 (D) OTHER INFORMATION:/label= Table 3  
3272  
3273 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:67 :  
3274  
3275 GGAGCTAGGATAGCTCAAT  
3276  
3277

# SEQUENCE VERIFICATION REPORT

## PATENT APPLICATION US/08/836,734

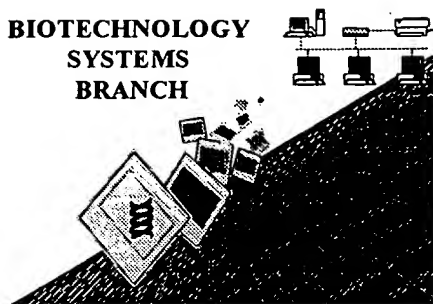
DATE: 06/16/98  
TIME: 13:37:41

INPUT SET: S26725.raw

Line	Error	Original Text
48	Entered (3018) and Calc. Seq. Length (0) differ	(A) LENGTH: 3018 base pairs
167	Entered (11451) and Calc. Seq. Length (0) differ	(A) LENGTH: 11451 base pairs
567	Entered (1834) and Calc. Seq. Length (0) differ	(A) LENGTH: 1834 base pairs
645	Entered (14664) and Calc. Seq. Length (0) differ	(A) LENGTH: 14664 base pairs
1152	Entered (5149) and Calc. Seq. Length (0) differ	(A) LENGTH: 5149 base pairs
1165	Unknown or Misplaced Identifier	(B) POSITION : 1303..3764
1170	Unknown or Misplaced Identifier	(B) POSITION : 1631
1176	Unknown or Misplaced Identifier	(B) POSITION : 1848
1182	Unknown or Misplaced Identifier	(B) POSITION : 1853
1188	Unknown or Misplaced Identifier	(B) POSITION : 2004
1194	Unknown or Misplaced Identifier	(B) POSITION : 2248
1200	Unknown or Misplaced Identifier	(B) POSITION : 2364
1206	Unknown or Misplaced Identifier	(B) POSITION : 2382
1212	Unknown or Misplaced Identifier	(B) POSITION : 2771
1218	Unknown or Misplaced Identifier	(B) POSITION : 3018
1224	Unknown or Misplaced Identifier	(B) POSITION : 3372..3373
1230	Unknown or Misplaced Identifier	(B) POSITION : 3533
1236	Unknown or Misplaced Identifier	(B) POSITION : 3609
1242	Unknown or Misplaced Identifier	(B) POSITION : 3616..3619
1248	Unknown or Misplaced Identifier	(B) POSITION : 3665..3666
2235	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2253	Entered (21) and Calc. Seq. Length (0) differ	(A) LENGTH: 21 base pairs
2271	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2289	Entered (21) and Calc. Seq. Length (0) differ	(A) LENGTH: 21 base pairs
2307	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2325	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2343	Entered (22) and Calc. Seq. Length (0) differ	(A) LENGTH: 22 base pairs
2362	Entered (19) and Calc. Seq. Length (0) differ	(A) LENGTH: 19 base pairs
2380	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2398	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2416	Entered (22) and Calc. Seq. Length (0) differ	(A) LENGTH: 22 base pairs
2434	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2452	Entered (22) and Calc. Seq. Length (0) differ	(A) LENGTH: 22 base pairs
2470	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2488	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2506	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2524	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2542	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2560	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2578	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2596	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2614	Entered (21) and Calc. Seq. Length (0) differ	(A) LENGTH: 21 base pairs
2632	Entered (21) and Calc. Seq. Length (0) differ	(A) LENGTH: 21 base pairs
2650	Entered (21) and Calc. Seq. Length (0) differ	(A) LENGTH: 21 base pairs
2668	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2686	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2704	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2722	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs

**SEQUENCE VERIFICATION REPORT**  
**PATENT APPLICATION US/08/836,734**DATE: 06/16/98  
TIME: 13:37:42*INPUT SET: S26725.raw*

Line	Error	Original Text
2740	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2758	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2776	Entered (22) and Calc. Seq. Length (0) differ	(A) LENGTH: 22 base pairs
2794	Entered (19) and Calc. Seq. Length (0) differ	(A) LENGTH: 19 base pairs
2812	Entered (21) and Calc. Seq. Length (0) differ	(A) LENGTH: 21 base pairs
2830	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2848	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2866	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2884	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2902	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2920	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2938	Entered (22) and Calc. Seq. Length (0) differ	(A) LENGTH: 22 base pairs
2956	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2974	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
2992	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
3010	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
3028	Entered (21) and Calc. Seq. Length (0) differ	(A) LENGTH: 21 base pairs
3046	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
3064	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
3082	Entered (19) and Calc. Seq. Length (0) differ	(A) LENGTH: 19 base pairs
3100	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
3118	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
3136	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
3154	Entered (22) and Calc. Seq. Length (0) differ	(A) LENGTH: 22 base pairs
3172	Entered (21) and Calc. Seq. Length (0) differ	(A) LENGTH: 21 base pairs
3190	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
3208	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
3226	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
3244	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
3262	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs



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